## STIC Database Tracking Number: 127078

TO: Terra Gibbs

Location: REM-2D10/2C18

Art Unit: 1635

Tuesday, July 13, 2004

Case Serial Number: 10/000213

From: Paul Schulwitz

**Location: Biotech-Chem Library** 

**REM-1A65** 

Phone: (571)272-2527

paul.schulwitz@uspto.gov

## Search Notes

Examiner Gibbs,

See attached results.

If you have any questions about this search feel free to contact me at any time.

Thank you for using STIC search services!

Paul Schulwitz Technical Information Specialist STIC Biotech/Chem Library (571)272-2527



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41 03H

## Schulwitz, Paul

From:

Gibbs, Terra

Sent:

Thursday, July 01, 2004 1:53 PM

To:

Schulwitz, Paul

Subject:

Sequence search request...

Hi David,

I have another request for a score over length search:

I need a length limited nucleotide sequence search of nucleobases 1710 through 1757 of SEQ ID NO:3 in USSN 10/000,213, where the returns are rank ordered based on the score over length/ratio as we've discussed. I need the lengths limited to hits between 8 and 80 nucleotides, and I'll take as many hits as you can import into excel (64,000?), and alignments for anything above .75 on the above ratio. Hope this is clear, please call me if it's not. I also need the interference databases searched.

Terra Cotta Gibbs, Ph.D. Art Unit 1635 Remsen Building 2D10 571-272-0758

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GenCore version 5.1.6 Copyright (c) 1993 - 2004 Compugen Ltd.

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Result
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ACCESSION: AR075067
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ACCESSION: AR828439
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Query Match Best Local Matches 1	LOCUS DEFINITION ACCESSION VERSION VERSION KEYWORDS SOURCE ORGANISM REFERENCE AUTHORS TITLE JOURNAL FEATURES BOURCE	Query Match Best Local Matches 1 Matches 1714  Qy 1714  Db 18  RESULT 2	KEYWORDS SOURCE ORGANISM REFERENCE AUTHORS TITLE JOURNAL FEATURES SOURCE	RESULT 1 AR075067/c LOCUS DEFINITION ACCESSION VERSION	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
l Similarity L6; Conserva	AR141885  Sequence 27 from patent US 614719 AR141885  AR141885.1 GI:15101401  Unknown. Unclassified. 1 (bases 1 to 18) Gimeno, C.J. and Errada, P.R. Tub interactor (TI) polypeptides Patent: US 6147192-A 27 14-NOV-20  Location/Qualifiers 118 /organism="unknown" /mol_type="unassigned_DN	h himilarity 16; Conserva 4 GCTGACTGATG                           8 GCTGACTGACG	Unknown. Unknown. Unknown. Unclassified. 1 (Gases 1 t Gimeno,C.J. a Genes encodin Patent: US 59 Patent: US 70 Patent) Patent (Jorg /mol.)	AR075067 Sequence 27 f AR075067 AR075067.1 G	7 7 7 8 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9
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Sequence 67 from Patent WO9517507.
A45190
          unidentified unclassified.
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Sequence 5840 from patent
AR328438
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AR336915
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Sequence
AR202833
                                                     1 (bases 1 to 16)

Brysch, W., Schlingensiepen, K.-H., Schlingensiepen, R. and Schlingensiepen, G.-F.
Antisense nucleic acids for the prevention and treatment disorders in which expression of c-erbB plays a role Patent: US 6365345-A 67 02-APR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 16)
Brysch, W. and Schlingensiepen, K.
ANTISENSE OLIGONUCLEOTIDE PREPARATION METHOD
PATENT: 09833904-A 1099 06-AUG-1998;
BIOGNOSTIK GES (DE); BRYSCH WOLFGANG (DE)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Brysch, W., Schlingensiepen, K., Schlingensiepen, R. and schlingensiepen, G.
ANTISENSE NUCLEIC ACIDS FOR THE PREVENTION AND TREATMENT DISORDERS IN WHICH EXPRESSION OF C-erbB PLAYS A ROLE Patent: WO 9517507-A 67 29-JUN-1995;
BIOGNOSTIK GES (DE)
Other publication, AU 1313095 950710.
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                                                                                                                                       Unclassified.
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nilarity 87.5%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                              /organism="unidentified"
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/db_xref="taxon:32644"
/organism≃"unknown"
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                                         Location/Qualifiers
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AR328439
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ACTIVE
AN antisense oligonucleotide preparation method
Patent: JF 2001511000-A 1099 07-AUG-2001;
BICANOSTIK GESELLSCHAFT FUR BICMOLEKULARE DIAGNOSTIK MBH
OS Unknown
PN JP 2001511000-A/1099
PD 07-AUG-2001
PF 30-JAN-1998 JP 1998532533
PF 31-JAN-1998 JP 1998532533
PF 31-JAN-1997 EP 97101577
PC C12N15/11
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Query Match 26. Best Local Similarity 87.9 Matches 14; Conservative
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BD066464
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 5841 from patent AR328439
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Pavco, P., McSwiggen, J.A.,
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                                                                                                                              ocation/Qualifiers
                                           /organism="unidentified"
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Location/Qualifiers
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/mol_type="unassigned RNA"
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Score 12.8; DB Pred. No. 6.9; 0; Mismatches
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KEYWORDS
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AX673162
                                                                                                                                                                                                     Telerman,A., Amson,R. and Tuijnder,M.
Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or virus resistance and their use as
                                                                                                                                                                                                                                                        Homo sapiens (human)
Homo sapiens
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent: WO 03004526-A 1607 16-JAN-2003;
Molecular Engines Laboratories (FR)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Telerman,A., Amson,R. and Tuijnder,M. Sequences involved in phenomena of tumour reversion, apoptosis and/or resistance to
                                                                                                                                                                  Patent: WO 03025175-A 5293 27-MAR-2003;
Molecular Engines Laboratories (FR)
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Mammalia; Eutheria; Primates;
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                                                                                                 /organism="Homo sapiens"
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/mol_type="unassigned DNA"
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87.5%;
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                                                            Score 12.8;
Pred. No. 7
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RESULT 12

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AR055874/c
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AR009449
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             AUTHORS
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                                                                                                                                                                                                                                                                                                           Unclassified.

1 (bases 1 to 15)

White,M.B. and Sadzewicz,L.K.

Susceptibility mutation for breast and
Patent: US 5756294-A 5 26-MAX-1998;

Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                           AR009449
Sequence
AR009449
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AR055874
Unclassified.
1 (bases 1 to 15)
Grimm,S., Stinchcomb,D.T.,
Draper,K.G.
                                                 Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Telerman,A., Amson,R. and Tuijnder,M.
Sequences involved in phenomena of tumour suppression,
Sequences and/or resistance to viruses and of
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Sequence 2652 from Patent
AX737062
                                                                                    AR055874.1
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Patent: WO 03025177-A 2652 27-MAR-2003;
                                                              Unknown
                                                                                                                       AR055874
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Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                              /organism="unknown"
/mol_type="unassigned
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                                                                                    GI:5981451
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86.7%; Pred. No. 10
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            McSwiggen, J.,
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             Sullivan,S.
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AX632925
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                                                                                                                                    Stinchcomb, D.T., Dudycz, L.W., Chowrira, B., Grimm, S., Direnzo, A., Karpeisky, A., Draper, K.G., Kisich, K., Metulic-Adamic, J., Mcswiggen, J.A., Modak, A., Pavco, P., Beigelman, L., Sullivan, S.M., Sweedler, D., Thompson, J.D., Tracz, D., Usman, N., Wincott, F.E. and
                                                                                                                                                                                                                       unidentified unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ribozyme treatment of diseases or conditions related to levels intercellular adhesion molecule-1 (ICAM-1) Patent: US 6132967-A 78 17-OCT-2000;
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Patent: US 5837542-A 78 17-NOV-1998;
Location/Qualifiers
                                                    Patent: EP 1260586-A 64 27-NOV-2002; RIBOZYME PHARMACEUTICALS, INC. (US) Location/Qualifiers
                                                                                                genes
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                                                                                                          Method and reagent for inhibiting the
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/organism="unidentified"
/mol_type="unassigned RNA"
/db_xref="taxon:32644"
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PR 27-APR-1998 US 60/083217,18-SEP-1998 US 60/100842 PR 27-APR-1999 US 09/257608,23-MAR-1999 US 09/274553 PI LAWRENCE BLATT,JAMES A MCSWIGGEN,ELISABETH ROBERTS,PAMELA A PI PAVCO, DENNIS MACEJAK PC C12N9/00,A61K31/7105,A61K38/21,A61K40/^-
PC A61K37/66, PC C12N15/00
CC Enverse
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M Unknown.

Unclassified.

E 1 (bases 1 to 15)

E 1 (bases 1, Kinzler, K.W., Zhang, L. and (S Vogelstein, B., Kinzler in normal and caprent: US 6333152-A 601 25-DEC-2001;
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/mol_type="genomic RNA"
/db_xref="taxon:32644"
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Sequence 8 :
AR013953
AR013953.1
 AX624126
                                                                                                                                                          Nucleic acid encoding calf intestinal alkaline phosphatase Patent: US 5707853-A 8 13-JAN-1998;
Location/Qualifiers
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Millan, J.L.
                                                                                                                                                                                                                    Unknown
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179681
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Millan,J.L.
Millan,J.L.
Recombinant calf intestinal alkaline
Patent: US 5773226-A 8 30-JUN-1998;
                                                                                                                                                                                                           Unclassified.
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                                                                                                                             /organism="unknown"
/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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Pred. No. 16;
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Pred. No. 12;
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                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                     Sequence 5681 from AX628640
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Sequence 2602 from Patent
AX625561
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1167 from Patent W002053774. AX624126
Method for determining homeostasis of the skin
                 Petersohn, D.,
                                                                        Homo sapiens
                                                                                         Homo sapiens (human)
                                                                                                                       AX628640.1 GI:28456678
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                 Conradt, M.
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100.0%; Pred. No. 18;
tive 0; Mismatches
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                 and Hofmann, K.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Henkel Kommanditgesellschaft auf Aktien
Location/Qualifiers
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Location/Qualifiers
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synthetic construct
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Synthetic EcoRI adaptor.
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M Unknown.
M Unknown.
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E 1 (bases 1 to 13)
RS McSwiggen.J.A.
Assay for Ribozyme target site
Patent: US 5525468-A 17 11-JUN-1996;
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Sequence 244
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Sequence 10 from patent US 5525468.
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121833.1 GI:1602187
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McSwiggen, J.A.
Assay for Ribozyme target site
Patent: US 5525488-A 10 11-UN-1996;
Location/Qualifiers
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Wang, C.-G. and Hepburn, A.G.
Genetic sequence assay using DNA triple
Patent: US 5861244-A 244 19-JAN-1999;
Location/Qualifiers
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244 from patent US 5861244.
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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Sequence
AR310639
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RIBOZYME PHARMACEUTICALS, INC. (US)
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AX711140
AX711140.1 GI:29787521
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Dervan, P.B., Wurtz, N. and Chang, A.
Polyamide-alkylator conjugates and related products and method
Patent: US 6559125-A 3 06-MAY-2003;
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                                                                                    /organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
                                                                          /note="Nucleic Acid"
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/mol_type="genomic DNA"
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Sequence 2700 from Patent
AX625659
             Homo sapiens (human)
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AX470508
AX470508.1 GI:22205633
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                     AX625659.1
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Method for determining skin stress or skin ageing in vitro
Patent: WO 02053773-A 85 11-JUL-2002;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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AX626369
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AX625973
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                                                                                                                                                                                                                                      Sequence 3410 from Patent AX626369
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/organism="Homo sapiens"
/mol_type="unassigned DNA"
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Sequence 5645 from Patent
AX628604
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                                /organism="Homo sapiens"
/mol_type="unassigned DNA"
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Sequence 8400 from Patent
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       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 7416 from Patent
AX630375
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Patent: WO 02053774-A 7416 11-JUL-2002;
Henkel Kommanditgesellschaft auf Aktien (DE)
Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Analysis of host cell shut-off
Patent: EP 0564801-A 16 13-OCT-1993;
BOEHRINGER INGELHEIM INT (DE)
Other publication DE 4206769 930909
Other publication JP 6197799 940719
Other publication CA 2090834 930905
Other publication DE 4217929 931202.
                                                                                                                                                12 bp DNA Short-chain oligonucleotide for inhibiting BD248202 BD248202.1 GI:33057972 JP 2002524038-A/21.
                                                                                                                                                                                                                                                                                         11 TTCATCCATTC 1
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artificial sequences.

1 (bases 1 to 12)

Uhlmann,E., Peyman,A., Bitonti,A. and Woessner,R.
Short-chain oligonuclectide for inhibiting VEGF e
Patent: JP 2002524038-A 21 06-AUG-2002;

AVENTIS PHARMA DEUTSCHLAND GMBH
OS Artificial Sequence
PN JP 2002524038-A/21

PD 06-AUG-2002
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A36017
                                                                                                                            synthetic construct
synthetic construct
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Sommergruber, W.D., Auer, H.,
Kuechler, E.P., Kowalski, H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       unidentified unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Petersohn,D., Conradt,M. and Hofmann,K. Method for determining homeostasis of the skin Patent: WO 02053774-A 8400 11-JUL-2002; Henkel Kommanditgesellschaft auf Aktien (DE) Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A36017
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                                                                                                                                                                                                                                                                                                                                                                                                       /organism="unidentified"
/mol_type="unassigned DNA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                              19.6%;
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1 (bases 1 to 12)

1 (bases 1 to 12)

1 (yamanishi, K., Yamamoto, T. and Mori, H. ANALYSIS OF HUMAN HERPES VIRUS 6 TYPE @(3754/24)HHV-6) DNA AND DISCRIMINATION OF SUB-TYPE

1 (Patent: JP 1994133799-A 10 17-MAY-1994;

1 (C)

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                                                1744 AAATGCATCCA 1754
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E07501
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Location/Qualifiers
                                                                                                                                                                                                    /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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/mol_type="genomic DNA"
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Matches 10; Conserv
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                      1730 GGAACAGACAG 1740
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OS None
OC Artificial sequences
PN JP 1994133799-A/25
PD 17-MAY-1994
PF 27-OCT-1992 JP 19923
PI YAMANISHI KOICHI, YAA
C1201/68,C1201/68,C12N15/
CC Strandedness Single
CC topology: Linear;
CC hypothetical: No;
CC anti-sense: Yes;
FH Key Loca
FT Source 1.
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12
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Baxter-Lowe,L.Ann.
Method for HLA Typing
Patent: US 5545536-A 11 13-AUG-1996;
Location/Qualifiers
                                                                                                                                                                                                                                                    Sequence 11 from patent US 124583 124583.1 GI:1604453
                                                                                                                                                                                                               Unknown
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Yamanishi,K., Yamamoto,T. and Mori,H.
Yamanishi,K., Yamamoto,T. and Mori,H.
ANALYSIS OF HUMAN HERPES VIRUS 6 TYPE @(3754/24)HHV-6)
DISCRIMINATION OF SUB-TYPE
Patent: JP 1994133799-A 25 17-MAY-1994;
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27-OCT-1992 JP 1992311416
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JP 1994133799-A/25
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                                                                                               /organism≃"unknown"
/mol_type="unassigned
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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                                                                                                                                                                         Telerman,A., Amson,R. and Tuijnder,M.
Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or virus resistance and their use as
                                                                                                                                                                                                                                                                                                                               AX733659 17 bp DNA Sequence 5293 from Patent WO03025175. AX733659
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OS Artificial Sequence
PN JP 2001275678-A/3
PD 09-OCT-2001
PF 31-MAR-2000 JP 2000096306
PI TOSHIHIKO KISHIMOTO, SHINICHIRO NIWA, YUKO MORI, SACHIYO
MIMAKI, REI FUKUSHIMA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     regulatory protein to oligoDNA
Patent: JP 2001275678-A 3 09-OCT-2001;
SUMITOMO ELECTRIC INDUSTRIES LTD
                                                                                                                                  Patent: WO 03025175-A 5293 27-MAR-2003;
Molecular Engines Laboratories (FR)
                                                                                                                                                                                                                                                                   Homo sapiens (human)
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1 (bases 1 to 12)

1 (bases 1, Niwa, S., Mori, Y., Kishimoto, T., Niwa, S.,
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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JP 2001275678-A/3.
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              Similarity
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                                                       /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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              83.3%;
                             18.3%;
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Pred. No. 23;
            Score 8.8; D
Pred. No. 31;
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Mismatches
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RESULT 50
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Best Local Similarity 88.9%;
Matches 8; Conservative
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Best Local Similarity 81.8%;
Matches 9; Conservative
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Synthetic EcoRI adaptor.
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A25126.1 GI:1247054
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synthetic construct synthetic construct
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Vogelstein,B., Kinzler,K.W., Zhang,L. and Zhou,W.
Gene expression profiles in normal and cancer cells
Patent: US 6333152-A 601 25-DEC-2001;
Location/Qualifiers
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Sequence 601 from patent US 6333152.
AR180533
AR180533.1 GI:20222566
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Method for determining homeostasis of the skin
Patent: WO 02053774-A 2602 11-JUL-2002;
Henkel Kommanditgesellschaft auf Aktien (DE)
Location/Qualifiers
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Unclassified.
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Pred. No. 47;
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/organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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75.0%;
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Pred. No. 49;
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                                                                                               ALIGNMENTS
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ADB99916;

ID ADB99916;

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XC ADB99916;

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XX Vitamin D

XX Cytostatic

KW vitamin D

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Best Local S
Matches 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cytostatic; gene therapy; antisense oligonucleotide; human; vitamin D nuclear receptor; cancer; developmental disorder; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                      Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense oligonucleotides for modulating vitamin D nuclear receptor gene expression, particularly useful for treating or preventing cancer odevelopmental disorder, or as diagnostics or research reagents.
                                                                                               04-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                    BP; 3 A; 8 C; 2 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                          Conservative
                                                                                               (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cytidine residues are 5-methylcytidines"
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/mod_base= OTHER
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                                                                                                                                                            DNA;
                                                                                             entry)
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Pred. No.
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RESULT 1 ADB99915/c ID ADB999, XX

ADB99915 standard; DNA; 20

ВP

Cytostatic; gene therapy; antisense oligonucleotide; human; vitamin D nuclear receptor; cancer; developmental disorder; phosphorothioate; ss.

Vitamin D nuclear receptor antisense oligonucleotide, SEQ ID

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Matches 20
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                                                                                                                                                                                            Vitamin D nuclear receptor antisense oligonucleotide,
                                                                                                                                                                                                                                                                                   ADB99914 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 1
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                                                                                                        Synthetic
                                                                                                                                   phosphorothioate; ss.
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                                                                                                                                                                                                                                                      ADB99914;
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                                                                                                                                                     vitamin D
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                                                           modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
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                                                                                                                                                    c; gene therapy;
nuclear recepto:
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                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                       (first entry)
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/mod_base= OTHER /mote= "This oligonucleotide has a phosphorothioate backbone and 2-'methyoxyethyl (2'-MOE) wings at the
                                                                         Location/Qualifiers
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                                                                                                                                                 therapy; antisense receptor; cancer;
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                                                                                                                                                  oligonucleotide; human; developmental disorder;
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Matches 20
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                                                                                                                                                                                                        Cytostatic; gene therapy; antisense oligonucleotide; human; vitamin D nuclear receptor; cancer; developmental disorder; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense oligonucleotides for modulating vitamin D nuclear gene expression, particularly useful for treating or preventing developmental disorder, or as diagnostics or research reagents.
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                                    22-MAY-2003.
                                                         WO2003041657-A2
                                                                                                                                                 modified_base
                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                       Vitamin D nuclear receptor antisense oligonucleotide,
                                                                                                                                                                                                                                                                                                      ADB99917;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-NOV-2002; 2002WO-US036692.
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                              (first entry)
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/mod= "This oligonucleotide has a phosphorothioate
backbone and 2-'methyoxyethyl (2'-MOE) wings at the
and 3' ends, which are 5 nucleotides in length. Also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and 3' ends, which are 5 nucleotides in length. Also cytidine residues are 5-methylcytidines"
                                                                                                                                    /*tag=
                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                             DNA;
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Pred. No.
                                                                                are 5-methylcytidines"
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13-NOV-2002; 2002WO-US036692.

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                                                       Disclosure; Page 85; 120pp; English.
                                                                                                                  Tub interactor genes - used to develop products for the treatment obesity, cachexia, anorexia nervosa or related disorders e.g. dial
                                                                                                                                                                                                                                                            Gimeno CJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antisense; tub interactor; treatment; obesity; cachexia; anorexia nervoa; diabetes; cell cycle progression; apoptosis; neurodegenerative diease; Alzheimer's disease; drug screening; Parkinson's disease; Huntington's chorea; detection; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense oligonucleotides for modulating vitamin D nuclear receptor gene expression, particularly useful for treating or preventing cancer or developmental disorder, or as diagnostics or research reagents.
                                                                                                                                                                                                                                                                                                                                                                         17-SEP-1996;
21-JUL-1997;
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                                                                                                                                                                                                                                                                                                                    (MILL-) MILLENNIUM PHARM INC
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97US-00897340.
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100.0%; r.
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                                                                                    Oligonucleotides, useful for the modulation of Smad6 expression in the treatment or prophylaxis of e.g. cardiovascular disease, are targeted nucleic acid molecule encoding Smad6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 18 BP; 3 A; 8 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                        WPI; 2002-394345/42.
                                                                                                                                                                                                                  Monia BP,
                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                  04-OCT-2000; 2000US-00679298.
                                                                                                                                                                                                                                                                                                                                          01-OCT-2001; 2001WO-US030645.
                                                                                                                                                                                                                                                                                                                                                                                         11-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200228878-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human Smad6 antisense oligonucleotide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-AUG-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1714 GCTGACTGATGTTGAGGG 1731
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Smad6 protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
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                                                                                                                                                                                                                Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cancer;
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/mod_bas
15. .18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag= b
/note= "2'methoxyethyl nucleotides"
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== "OTHER =
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       c
"2'methoxyethyl nucleotides"
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Pred. No. 23
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RESULT 7
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; zinzyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nucleic acid molecule encoding human Smad6 protein, which specifically hybridises with the nucleic acid and inhibits its expression. Antisense compounds of the invention are used for inhibiting the expression of smad6 in cells and tissues in the treatment of a disease or condition associated with Smad6 such as cardiovascular disease, cancer, infection and inflammation. They are also useful in the diagnostics, as research reagents, in kits and in antisense therapy. The present sequence is an
                                                                                                                                                                                                  (RIBO-)
(BLAT/)
(MACE/)
(MCSW/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 BP; 3 A; 7 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                              Blatt
                                                                                                                                                                                                                                                                                                                  26-MAR-2001; 2001US-00817879.
                                                                                                                                                                                                                                                                                                                                           26-MAR-2002; 2002WO-US009187.
                                                                                                                                                                                                                                                                                                                                                                    17-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                            WO200281494-A1
                                                                                                                                                                                                                                                                                                                                                                                                                    Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HCV minus strand DNAzyme substrate sequence #749.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-SEP-2003
Claim 1; Page 288; 387pp; English
                                   Novel compound useful for hepatocellular carcinoma,
                                                                           WPI; 2003-229207/22.
                                                                                                                                                                                                                                                                                         08-JUN-2001;
                                                                                                                                                                                                                                                                                                     08-JUN-2001;
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                                                                                                 ~
                                                                                                                                                                        BLATT L.

MACEJAK D.

MCSWIGGEN J.

MORRISSEY D.

PAVCO P.
                                                                                                                                     LEE P.
DRAPER K.
ROBERTS E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                       RIBOZYME PHARM
                                                                                                 Macejak D,
Roberts E;
                                                                                                                                                                                                                                                                                                                                                                                                                                             antiinflammatory; substrate; ss.
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2001US-0296876P.
2001US-0335059P.
2001US-0337055P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entry)
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                                                                                                              Mcswiggen J,
                                                                                                                                                                                                                                         INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1738
                                      treating cirrhosis, or condition associ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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Pred. No. 2
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                                                                                                             Morrissey
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                                      associated
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                                      liver failure,
ated with hepat
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                                                                                                              Pavco
                                    hepatitis C
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                                                                                                              Lee
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RESULT 8
ABNO8350/c
ID ABNO8350/c
ID ABNO835
XX ABNO83
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XX HUMAN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ouery Match
Best Local Similarity
Matches 11; Conserv
                                                                                                                                      26-MAY-2000; 2000US-0237456P.
21-SEP-2000; 2000US-0236359P.
27-SEP-2000; 2000US-0236359P.
04-CCT-2000; 2000GB-00024263
30-JAN-2001; 2001WO-US000661.
30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000668.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; hear muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABN08350 standard; DNA; 17
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                                                                                        (AEOM-) AEOMICA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAY-2001; 2001WO-US016981.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-MAY-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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                              SG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28.7%;
                              Hanzel
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Pred. No. 30
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                              봈
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                                 Rank
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                                 Chen
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                                 Ξ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ħ
                                 Shannon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NO:8342.
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                                 M
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Penn

2002-179446/23

Disclosure; SEQ ID NO 8342; 214pp; English

w polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, as specific biomolecule capture probes for surface-enhanced laser sorption ionization, comprises human myosin-like protein hGDMLP-1.

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RESULT 9
ABNO8352/c
ID ABNO8352;
XX
AC ABNO8352;
XX
DT 29-MAY-2002 (first entry)
XX
Human GDMLP-1 17-mer scanning
XX

DE Human GDMLP-1 17-mer scanning
XX

Muscle tal muscle disorder; amp
XX
Homo sapiens.
XX
PN W0200192524-A2.
XX
PN W0200192524-A2.
XX
PN W72001; 2001W0-US016981.
XX
PN W0200192524-A2.
XX
PN W
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The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from MIPO as fer wine int/puh/mblished not sequence.
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence
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         ; 2000US-0207456P.
; 2000US-023458PP.
; 2000US-0236359P.
; 2000GB-00024263.
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; 2001WO-US000663.
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Pred. No.
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CC expressing the proteins. The hCDMLP-1 proteins or polypeptides may be cused as immunogens to raise antibodies that specifically recognise hGDMLP proteins, as standards in assays used to determine the concentration and/or amount specifically of hCDMLP proteins, as specifically of hCDMLP proteins, as specific blomolecule capture probes for surface-enhanced laser desorption ionisation, as the respective probes for surface-enhanced laser desorption ionisation, as complement in patients having specific deficiency in hCDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hCDMLP-1 may be used for diagnosing a clisorder associated with the expression of hCDMLP-1 in particular heart and skeletal muscle disorders. hCDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the present sequence in the exemplification of the present invention. N.B.

The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO as for which into the present directly from WIPO.
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30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000670.
05-FEB-2001; 2001US-0266860P.
                                                                                                                                                                                                                                                                                                                                                                                            nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         rotein 1 (hGDMLP-1). The protein and polynucleotide sequences of can be used in gene therapy and vaccine production. The hGDMLP-1
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                                      ftp.wipo.int/pub/published_pct_sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sent invention
1 (hGDMLP-1).
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Query Match Best Local Matches 1713 TGCTGACTGATGT 1725 Similarity Conservative 27.1%; 0 Score 13; ; Pred. No. Mismatches DB 1; 39; 0; Length 17 Indels

<u>,,</u>

0;

Sequence 17

BP; 6 A; 4 C; 5 G; 2 T; 0 U; 0 Other;

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ABNO8351/
ID ABNO
XX
AC ABNO
XX
AC ABNO
XX
UP Huma
XX
Homc
XX RESULT 10 ABN08351; ABN08351 standard; ີດ DNA;

Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:8343.

29-MAY-2002

(first entry)

Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapv; vaccine: myosin; chromosome l muscle disorder; a 22; gene therapy; vaccine; heart disease; amplicon; screening; ss. hGDMLP-1; heart;

Homo sapiens.

WO200192524-A2

25-MAY-2001;

2001WO-US016981

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RESULT 11
ABN08349/c
ID ABN083
XX
AC ABN083
XX
DT 29-MAY
XX
DE Human 1
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                                                                                                                                                                                                                                                                                                                         The present invention describes a human genome-derived myosin-like CC protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 cC nucleic acids can be used as probes to detect, characterise and quantify cc provide initial substrates for the recombinant engineering of hGDMLP-1 cC protein variants having desired phenotypic improvements, and for cexpressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP cc and/or amount specifically of hGDMLP proteins, as standards in assays used to determine the concentration cc apture probes for surface-enhanced laser describion ionisation, as the action and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The concentration acc disorder associated with the expression of hGDMLP-1 may be used for diagnosing a cd disorder associated with the expression of hGDMLP-1, in particular heart cand skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence in the expresents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present invention. N.B. Cf hGDMLP-1 sequence in the exemplification of the present invention. N.B. considered the protein of the printed cc specification, but was obtained in electronic format directly from WIPO cc at ftp.wipo.int/pub/published_pct_sequence
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Best Local :
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30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
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21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence SEQ ID NO:8341
                               29-MAY-2002
                                                              ABN08349
                                                                                                ABN08349
                                                                                                                                                                                                                                                                                                     Sequence
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30-JAN-2001;
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                                                                                              standard; DNA; 17
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2000US-0234687P.
2000US-0236359P.
2000US-0236359P.
2001WO-US000661.
2001WO-US000662.
2001WO-US000664.
2001WO-US000665.
2001WO-US000665.
2001WO-US000666.
2001WO-US000666.
2001WO-US000666.
2001WO-US000666.
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                               (first entry)
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Pred. No.
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27-SEP-2000;
04-OCT-2000;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.
                                     05-FEB-2001;
                                                   30-JAN-2001;
30-JAN-2001;
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30-JAN-2001;
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21-SEP-2000;
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(AEOM-) AEOMICA INC.
                               2000GB-00024263

2001WO-US000661

2001WO-US000663

2001WO-US000664

2001WO-US000665

2001WO-US000666

2001WO-US000667

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      heart disease;
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New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1. WPI; 2002-179446/23. Penn SG,

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Υ,

Hanzel DK,

Rank DR,

Chen

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Shannon ME;

Disclosure; SEQ ID NO 8341; 214pp; English.

CC protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 cn used in gene therapy and vaccine production. The hGDMLP-1 cc nucleic acids can be used as probes to detect, characterise and quantify cc hGDMLP-1 nucleic acids in samples, as amplification substrates, to protein variants having desired phenotypic improvements, and for cc provide initial substrates for the recombinant engineering of hGDMLP-1 cc protein variants having desired phenotypic improvements, and for cc expressing the proteins. The hGDMLP-1 proteins or polypeptides may be cc. 1 proteins, as standards in assays used to determine the concentration cand/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as concentration, and in vaccines or for replacement therapy. The conduction, and in vaccines or for replacement therapy. The colorated sequences encoding hGDMLP-1 may be used for diagnosing a colorated with the expression of hGDMLP-1, in particular heart colorates associated with the expression of hGDMLP-1, in particular heart colorates associated with the expression of hGDMLP-1, in particular heart colorates associated with the expression of hGDMLP-1, in particular heart colorates associated with the expression of hGDMLP-1, in particular heart colorates associated with the expression of hGDMLP-1, in particular heart colorates associated associated with the expression of hGDMLP-1, in particular heart colorates associated associated with the expression of hGDMLP-1, in particular heart colorates associated with the expression of hGDMLP-1, in particular heart colorates associated with the expression of hGDMLP-1, in particular heart colorates associated with the expression of hGDMLP-1, in particular heart colorates associated with the screening of the present sequence associated with the screening of the present sequence associated with the screening of the present directly from the screening of the particular heart with the screening of the present sc The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of a ftp.wipo ification, on, but was obtained in electronic format directly int/pub/published\_pct\_sequence from WIPO

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Sequence 17
 BP; 7
 A; 5 C; 3 G; 2 T; 0 U; 0 Other;
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밁
                                 Query Match
Best Local S
Matches 13
            1713 TGCTGACTGATGT 1725
17 TGCTGACTGATGT 5
                                          Similarity
                                  Conservative
                                         27.1%;
                                  ۰,
                                  Score 13; DB 1;
Pred. No. 39;
0; Mismatches
                                  0
                                                  Length 17
                                   Indels
                                  0;
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The present invention describes a human genome-derived myosin-like protein 1 (hgDMLP-1). The protein and polynucleotide sequences of hgDMLP-1 CC l can be used in gene therapy and vaccine production. The hgDMLP-1 CC mucleic acids can be used as probes to detect, characterise and quantify CR hgDMLP-1 nucleic acids in samples, as amplification substrates, to CC provide initial substrates for the recombinant engineering of hgDMLP-1 CC protein variants having desired phenotypic improvements, and for CC expressing the proteins. The hgDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hgDMLP-1 cc used as immunogens to raise antibodies that specific biomolecule capture proteins, as standards in assays used to determine the concentration cc and/or amount specifically of hgDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as CC therapeutic supplement in patients having specific deficiency in hgDMLP-1 cc polynucleotide sequences encoding hgDMLP-1 may be used for diagnosing a CC disorder associated with the expression of hgDMLP-1, in particular heart CC and skeletal muscle disorders. hgDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hgDMLP-1 sequence in the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the printed CC specification, but was obtained in electronic format directly from WIPO
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ABN08353/c
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21-SEP-2000; 2000US-0234587P.
27-SEP-2000; 2000US-0235359P.
04-CCT-2000; 2000GB-00024263
30-JAN-2001; 2001WO-US000661.
30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
30-JAN-2001; 2001WO-US000669.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-DEC-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     muscle;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human GDMLP-1 17-mer scanning SEQ ID NO:5 sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABN08353 standard, DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-MAY-2001; 2001WO-US016981.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200192524-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                skeletal muscle
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABN08353;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ,
Y
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart;
; myosin; chromosome 22; gene therapy; vaccine; heart disease;
al muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AEOMICA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Υ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 8345; 214pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Penn SG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rank DR,
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RESULT 13
AAQ92724/5
AAQ92724/5
AC AAQ92724;

XX
AC AAQ92724;
XX
DT 13-FEB-1996 (first entry)
YX
C-erbB-2 antisense nucleic
XX
DE c-erbB-2 antisense nucleic
XX
DE c-erbB-2 antisense nucleic
XX
DE c-erbB-2 antisense nucleic
XX
Antisense nucleic acid; c-
XX
PN W09517507-A1.
XX
PN W09517507-A1.
XX
PD 29-JUN-1995.
XX
PD 29-JUN-1995.
XX
PD 29-JUN-1995.
XX
PT (BIOG-) BIOGNOSTIK GES BIO
XX
PT (BIOG-) BIOGNOSTIK GES
AN (BIOG-) BIOGNOSTIK
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                                                                                                       Query Match
Best Local S
Matches 14
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Best Local 9
                                                                                                                                                                                                                                                                                          The sequences given in AAQ92658-762 are antisense nucleic acids which hybridise with part of the mRNA and/or DNA encoding c-erbB-2. These antisense nucleic acids are able to inhibit the expression of the p185-erbB-2 protein tyrosine kinase activity and cell growth in a number of tumour cells including breast cancer cells. Untransformed normal fibroblasts are not growth inhibited by anti-c-erbB-2 antisense compounds suggesting that p185-erbB-2 plays a pathogenic role in the growth of the above mentioned tumours. These antisense oligonucleotides may be used in the prevention and treatment of neoplasms, immune diseases and/or diseases involving pathological angiogenesis when associated with c-erbB-2 expression. They may also be used to detect expression of the relevant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New anti:sense nucleic acid against C-erbB-2 neoplasms, immune disease and angiogenesis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 35; 55pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Antisense nucleic acid; c-erbB-2; inhibition; fibroblast; neoplasm; p185-erbB-2 protein tyrosine kinase; tumour; breast cancer; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   c-erbB-2 antisense nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17 BP; 7 A; 4 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       at ftp.wipo.int/pub/published_pct_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (BIOG-) BIOGNOSTIK GES BIOMOLEKULARE DIAGNOSTIK.
                            1723 TGTTGAGGGAACAGAC 1738
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1713 TGCTGACTGATGT 1725
16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         disease; angiogenesis; ss.
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                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Schlingensiepen K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ilarity 100.0%;
Conservative 0;
                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                93EP-00120710
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       94WO-EP004094
                                                                                                                                                                                                                    5 C;
                                                                                                                                 26.7%;
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                                                                                                                                                                                                                    G; 6
                                                                                                       0
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Pred. No.
                                                                                                                                 Score 12.8;
Pred. No. 40
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                                                                                                          Mismatches
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                                                                                                                                                                                                                    T; 0
                                                                                                                                                                                                                    Ç;
                                                                                                                                    40;
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                                                                                                                                                                                                                       0
                                                                                                                                                              DB
                                                                                                                                                                                                                       Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 - for treating or
also for diagnosis.
                                                                                                                                                              1; Length 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     preventing
                                                                                                          Gaps
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RESULT

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RESULT 15
ABT39656
ID ABT39
XX ABT39
XX ABT39
XX 12-JU
XX Cytos
KW Cytos
KW anti:
KW anti:
KW human
XX homo
XX Homo
XX WO20
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ID ACC52840 standard;
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Best Local S
Matches 14
                                                               Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; geantisense; sense; tumour; cell degeneration; cancer; Alzheimer's schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.
                                                                                                                                                                                                                                                                                                                                                                                                             with tumour suppression or regression, apoptosis or virus resistance. invention relates to these sequences or sequences having at least 80% identity to them, and polypeptides encoded by the sequences or polypeptides having 80% identity to the polypeptide sequences. The invention is used to diagnose or treat viral disease or disease characterized by development of tumour cells or cellular degeneration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New nucleic acid sequences associated with tumor suppression, rappoptosis or virus resistance are useful to diagnose and treat disease, development of tumor cells and cell degeneration.
                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human tumour suppressor sequence #1607
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                                        Homo sapiens
                                                                                                                                Tumour suppression related human
                                                                                                                                                            12-JUN-2003
                                                                                                                                                                                      ABT39656
                                                                                                                                                                                                                 ABT39656 standard; DNA; 17 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tuijnder M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-JUN-2001; 2001FR-00008139.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cellular degeneration.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  tumour regression; apoptosis; virus resistance; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ss; tumour suppressor; antitumour; cytostatic; tumour suppression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACC52840;
             WO2003025175-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (MOLE-)
                                                                                                                                                                                                                                                                                                        1737 ACAGGAGAAATGCATC 1752
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequence represents an isolated nucleic acid sequence associated
                                                                                                                                                                                                                                                                                16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Page 411; 798pp;
                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULAR ENGINES LAB SA.
                                                                                                                                                                                                                                                                                                                                                                                      BP; 1 A; 6 C; 2 G; 8 T; 0 U;
                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                               26.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Amson R;
                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                           Score 12.8;
                                                                                                                                                                                                                                                                                                                                               Pred.
                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                  fukutin oligo SEQ
                                                                                                                                                                                                                                                                                                                                               No.
                                                                                                                                                                                                                                                                                                                                                                                       0 Other;
                                                                                                                                                                                                                                                                                                                                                             DB 1;
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                                                                                                                                                                                                                                                                                                                                                           Length 17;
                                                                                                                                  ID No
                                                                                                                                                                                                                                                                                                                                    0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                regression,
                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                           ene chip;
disease;
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ACD59839/C
ID ACD598
XX AC

HCV DNAzyme substrate sequence #1529.

24-SEP-2003 ACD59839;

(first entry)

ACD59839 standard; RNA; 17

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Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; zinzyme

amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrh liver failure; hepatocellular carcinoma; hepatotropic; cytostati

Hepatitis

a

virus

virucide;

antiinflammatory;

substrate; ss.

cytostatic; cirrhosis; 밁 S

1731 GAACAGACAGGAGAAA 1746

1 GATCAGGCAGGAGAAA 16

Query Match Best Local S Matches 14

Similarity

26.7%;

Score 12.8; Pred. No. 4: Mismatches

BG ۲. 2

Length 17; Indels

0

Gaps

0

14;

Conservative

0

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The invention relates to a novel isolated 17 mer nucleic acid sequence, given in the specification, a sequence containing at least 15 consecutive conditions at least 17 mer sequence with, after optimal configurations at least 80 % identity to the 17 mer sequence with, after optimal configurations at least 80 % identity to the 17 mer sequence, a sequence that configuration of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, conditions of the invention are useful as probes and primers for detecting, component of a gene chip, in vitro as (anti)sense reagents, and for component of a gene chip, in vitro as (anti)sense reagents, and for component of recombinant polypeptides. Any of the nucleic acids, e.g. as one conjugations, vectors containing the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for component of pharmaceuticals for prevention and/or treatment of viral consess that are characterised by development of tumours or cell containing the expression of the 17 mer nucleic acids in containing the expression of the 17 mer nucleic acids in containing the polypeptides can also be used to generate antibodies, and both the polypeptide and antibodies are useful as components of these contains. The nucleic acid sequences of the invention can be used in gene therapy. This polyvuclentide sequence represents a tumour suppression
                                                                chips. The nuc-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-SEP-2002; 2002WO-IB004208
Sequence 17
                                            therapy. This polynucleotide sequence represents a tumour suppression related human fukutin oligonucleotide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 652; 720pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-SEP-2001; 2001FR-00011978
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MOLE-) MOLECULAR ENGINES LAB.
BP; 8 A; 2 C; 6 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Amson R,
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RESULT 17
ABF92692
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                                                                                                                                                                                                                                                                   The present invention relates to nucleic acid molecules which modulate CC the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense compounds in the present cacids such as hammerhead ribozymes, DNAzymes, CI inozymes, zinzymes, and G-cleaver ribozymes. Also disclosed care nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV CC DNA. The nucleic acids may be used to modulate the expression of HBV compounds and/or potential therapies directed against HBV, and compounds compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and disease states related to HBV and HCV infection, replication and gene caxpression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represents a substrate for one of the HCV invention or minus strand DNAzyme sequences disclosed in the present
                                                                                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-MAR-2001;
08-JUN-2001;
08-JUN-2001;
08-JUN-2001;
24-OCT-2001;
05-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Blatt L,
Draper K,
           Oligonucleotide SEQ ID NO 192689 for detecting SNP TSC0047415.
                                     22-FEB-2002
                                                                                        ABF92692 standard; DNA; 13
                                                                                                                                                                                                                                                                   Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; Page 261; 387pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel compound useful for hepatocellular carcinoma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-229207/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (RIBO-)
(BLAT/)
(MACE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-MAR-2002; 2002WO-US009187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200281494-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ROBE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (MCSW/)
                                                                                                                                                                            1715 CTGACTGATGTTGAGG 1730
                                                                                                                                                                                                               14;
                                                                                                                                                           17
                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DRAPER K.
ROBERTS E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MCSWIGGEN J.
MORRISSEY D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RIBOZYME PHARM INC. BLATT L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PAVCO P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MACEJAK D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LEE P.
                                                                                                                                                           CTGAGTGATGGTGAGG
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Roberts E;
                                                                                                                                                                                                                                                                   BP; 4 A;
                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2001US-00877478.
2001US-0296876P.
2001US-0335059P.
2001US-0337055P.
                                     (first entry)
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                                                                                                                                                                                                                           26.7%;
                                                                                                                                                                                                                                                                  8 С; 2 G; 0 Т; 3 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mcswiggen J,
                                                                                                                                                           N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            treating cirrhosis, liver failure, or condition associated with hepatitis C virus
                                                                                                                                                                                                              0;
                                                                                                                                                                                                                            Score 12.8;
Pred. No. 4:
                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Morrissey
                                                                                                                                                                                                                             42;
                                                                                                                                                                                                                                                                   0 Other;
                                                                                                                                                                                                                                         DB 1; Length 17;
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                                                                                                                                                                                                               Indels
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomers are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                    Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-657177/75.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-APR-2000; 2000DE-01019173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-APR-2001; 2001WO-IB000713.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200177384-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EPIGENOMICS
                                                                                                                                                                                                                                                                                   SEQ ID NO 192689; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                    Piepenbrock C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ĄG.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Berlin
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                                                                                                                                                                                                                                                                                                                                          typing, :
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S Query Match
Best Local Similarity
Matches 12; Conserv Sequence 13 BP; 2 A; 0 C; 6 G; 4 T; 0 U; 1 Other; 1720 TGATGTTGAGGG 1731 1 TGATGTTGAGGG 12 Conservative 25.0%; <u>.</u> Score 12; Pred. No. Mismatches DB 1; 0; Length 13 Indels 0 Gaps 0;

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RESULT 18
ABF92693/c
ID ABF926
XX ABF926
XX ABF926
XX Oligon
XX SNP; BS
KW SNP; BS
KW SPPtid
KW Centra
XX Peptid
KW Centra
XX Homo B
XX HOMO B
XX HOMO B
XX HOMO B
XX WO2001
XX O6-APR
XX O6-APR
XX O7-APR
                                                                                                                                                                                                                                                                                                                                                                           밁
                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                            18-OCT-2001.
                                                                                         WO200177384-A2.
                                                                                                                                                                                                          Oligonucleotide SEQ
                                                                                                                                                                                                                                         22-FEB-2002
                                                                                                                                                                                                                                                                       ABF92693;
                                                                                                                                                                                                                                                                                                 ABF92693 standard; DNA; 13
                                                                                                                     sapiens.
                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                            ID NO 192690 for
                                                                                                                                                                                                                                                                                                    ВÞ
                                                                                                                                                                                                          detecting
                                                                                                                                                                                                             SNP TSC0047415.
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07-APR-2000; 2000DE-01019173. 06-APR-2001; 2001WO-IB000713.

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RESULT 19
AAT51864/c
ID AAT518
XX AAT518
XX O9-MAR
DT 25-MAR
DT 09-MAR
XX U9-MAR
XX U10-MAR
XX Enzyma
KW Enzyma
KW Enzyma
KW transl
KW t
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Best Local S
Matches 12
  23-FEB-1994;
29-MAR-1994;
04-APR-1994;
07-APR-1994;
15-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation, ABC00010-ABC9989, ABF00010-ABF99989, ABF00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. MOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                       intercellular adhesion molecule; rel A; tumour necrosis factor; TNF-alpha; respiratory syncytial virus; RSV; bcr-abl; oncogene; translocation; chronic myelogenous leukaemia; CML; cancer; philadelphia chromosome; inflammation; autoimmune disease; atherosclerosis; myocardial infarction; stroke; restenosis; transplant rejection; remmatoid arthritis; psoriasis; myocardial ischaemia; Kawasaki disease; septic shock; HIV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Enzymatic nucleic acid; ribozyme; trans cleavage; inhibition; gene expression; downregulation; interleukin-5; IL-5; ICAM-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-MAR-2003
09-MAR-1997
                                                                                                                                                                         23-FEB-1995;
                                                                                                                                                                                                                                 31-AUG-1995
                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT51864;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT51864 standard; RNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Olek A,
                                                                                                                                                                                                                                                                                    WO9523225-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                            immunodeficiency virus; acquired
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ICAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ch 25.0%; So 1 Similarity 100.0%; 1 12; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EPIGENOMICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hammerhead ribozyme target sequence (nt. position
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ID NO 192690; 29pp + Sequence Listing; German.
  94US-00201109.
94US-00218934.
94US-00222795.
94US-00224483.
94US-00227958.
                                                                                                                                                                            95WO-IB000156.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      å
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1731
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ζ.
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                                                                                                                                                                                                                                                                                                                                                                                                                               immune deficiency syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cytosine
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ARESULT 20
AAZ6422/C
ID AAZ642
XX AAZ642
XX AAZ642
XX BABET
XX BABET
XX Substr
XX Substr
XX Enzyma
XX autoim
XX autoim

28-MAR-2000

(first

entry)

for

hammerhead ribozyme which cleaves HCV

RNA

at

AAZ64202 standard;

RNA;

15 ₽₽

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Enzymatic nucleic acid; hammerhead ribozyme; virus replication; cleavage; cirrhosis; liver failure; hepatocellular carcinoma; interferon; cancer; autoimmune disease; ss.

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                                                                                                            Matches
                                                                                                                                   Query Match
Best Local Similarity
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23-SEP-1994;
23-SEP-1994;
23-SEP-1994;
28-SEP-1994;
03-OCT-1994;
11-OCT-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-AUG-1994;
17-AUG-1994;
19-AUG-1994;
                                                                                                                                                                                                                                                                    The present sequence represents a preferred target sequence for an enzymatic nucleic acid (i.e. a ribozyme) which cleaves ICAM-1 mRNA. Regions of the mRNA that do not form secondary folding structures and that contain potential hammerhead and hairpin ribozyme cleavage sites were identified by computer analysis. Ribozymes directed against these mRNA sequences were designed and synthesised with modifications that improve their nuclease resistance. The ribozymes cleave the ICAM-1 target sequences and thereby inhibit ICAM-1 expression, making them useful for reducing transplant rejection and alleviating symptoms in patients with rheumatoid arthritis, asthma and other inflammatory disorders. (Updated on 25-MAR-2003 to correct PI field.)
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Modak
                                                                                                                                                                                                                     Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ribozymes having modified bases and methods in inhibiting disease related genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1995-351090/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tracz
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15-AUG-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-APR-1994;
18-MAY-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (RIBO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2; Page 172; 407pp; English
15 GTCCAGGGAACAGAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RIBOZYME PHARM INC
                                                 GTTGAGGGAACAGAC 1738
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                                                                                                                                                                                                                     BP; 1 A; 5 C; 4 G; 0 T; 5 U; 0 Other;
                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       94US-00228041
94US-00245736
94US-00291932
94US-00291932
94US-00291620
94US-00293520
94US-00303039
94US-00311486
94US-00311749
94US-003114397
94US-003114397
94US-00311497
94US-00311497
94US-00311497
94US-0031193
94US-00337608
94US-00345516
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                                                                                                                                   24.6%;
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                                                                                                                                   Score 11.8;
Pred. No. 53;
                                                                                                            Mismatches
                                                                                                                                                             DB
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                                                                                                                                                             1; Length 15;
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ARESULT 21
AAF46891
ID AAF46
AC AAF46
AC AAF46
AC AAF46
XX 30-MA
XX 30-MA
XX Antis
KW Cytos
KW Cytos
KW Cytos
KW IGF L
K
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18-SEP-1998;
25-FEB-1999;
23-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence represents the preferred target sequence of an enzymatic nucleic acid, especially a hammerhead ribozyme, which cleaves the Hepatitis C virus (HCV) RNA sequence at the base position given in the descriptor line. The HCV sequence was screened for optimal ribozyme target sites using a computer folding algorithm and regions of the mRNA which did not form secondary folding structures and contained potential ribozyme cleavage sites were identified. Ribozymes were synthesised to target these sites and their activities optimised by either varying the length of the binding arms or by modification to prevent degradation by nucleases. The ribozymes of the invention inhibit gene expression and/or viral replication, and are used to treat diseases associated with Hepatitis C virus (HCV) infection, e.g. cirrhosis, liver failure and hepatocellular carcinoma. The ribozymes may be used in combination with
                                                                                                                                                                                                                       Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hepatocellular carcinoma. The ribosinterferon to treat HCV infection, diseases, and cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Blatt L,
                                                             Homo sapiens
                                                                                                                                           keratosi8; neoplasia; scleroderma; wart; skin c;
hyperneovascular condition; hyperplasia; kidney
                                                                                                                                                                                             growth factor mediated cell proliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                             30-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF46891 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 84; 123pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel ribozymes for the treatment of diseases and conditions related to hepatitis {\tt C} infection.
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                                                                                                                 neovascular condition of the retina;
                                                                                                                                                                                                                                                                                                                                                                      IGFBP3 oligonucleotide #311.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (RIBO-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 4 A; 7 C;
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                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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98US-0100842P.
99US-00257608.
99US-00274553.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Roberts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 11.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      u;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           53;
                                                                                                                                                                      n; ichthyosis; serborrhoea; ruba; skin cancer; sclerotic disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              infectious
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              autoimmune
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an CC antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 CC receptor, IGF binding protein [IGFBP]-2 or IGFBP3], which is capable of inhibiting or reducing growth factor mediated cell proliferation, CC inflammation and/or other disorders. The present sequence is an CC oligonucleotide which can be used to design the antisense CC oligonucleotides of the present invention (see AAF45151 and AAF45153-CC (1990). The method is useful for ameliorating the effects of psoriasis, CC (1990) is pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, CC (1990) is pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, CC (1990) is pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, CC (1990) is pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, CC (1990) is pityriasis, ruba, pilaris, cancers of the skin, a CC (1990) is pityriasis, ruba, pilaris, cancers of the skin, a CC (1990) is pityriasis, ruba, pilaris, cancers of the skin, a CC (1990) is pityriasis, ruba, pilaris, cancers of the skin, a CC (1990) is pityriasis, ruba, pilaris, cancers of the skin, a CC (1990) is pityriasis, ruba, pilaris, cancers of the psecular condition of the retina, ruba, pityriasis, ruba, pilaris, cancers of the psecular condition of the retina, ruba, pityriasis, ruba, pityriasis, ruba, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, ruba, pilaris, serborrhoea, keloids, ruba, pilaris, serbor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local (
                                                                                                                                                                                                                            Antisense therapy; antiproliferative; antinflammatory; antipsoriatic; cytostatic; dermatological; acardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1, pitryiasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhea; ruba; keratosis; neoplasia; scleroderma, wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15
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                                                                                                                                                                                                        neovascular condition of the retina; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                               IGFBP3 oligonucleotide #312.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAF46892 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 7; Page 46; 201pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1710 GGCTGCTGACTGATG 1724
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ر
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Werther GA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             other hyperplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99US-0140345P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24.6%;
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Pred. No. 5:
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21-JUN-2000; 2000WO-AU000693.

28-DEC-2000 WO200078341-A1

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cc antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 cc receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, cc inflammation and/or other disorders. The present sequence is an cc oligonucleotide which can be used to design the antisense oligonucleotide which can be used to design the antisense oligonucleotide of the present invention (see AAF45.51 and AAF45153-cc F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, cc ichthyosis, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, cc brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood cc vessels or any other hyperplasia
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                              Cytochrome P450; dioxin-inducible; glaucoma 3; CYPIB1; cytostatic; ophthalmological; gene therapy; polymorphism; breast cancer; ASO; primary congenital glaucoma; allele-specific oligonucleotide; prob
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                                                       WO200230951-A2
                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                            ASO probe #7 for detecting CYP1B1 gene polymorphisms.
                                                                                                                                                                                                                                                                                                                                                           16-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                        ABN81214 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 15 BP; 1 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 7; Page 46; 201pp; English
                 WPI; 2002-426265/45
                                                Han J,
                                                                                                           13-OCT-2000; 2000US-0240211P
                                                                                                                                          15-OCT-2001; 2001WO-US042726.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          present invention relates to a method for ameliorating the effects
                                                Kliem SE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                              GENAISSANCE PHARM INC
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                                                 Sanchis A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              53;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
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RESULT 24
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ID ABXO12
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XX ABXO12
XX ABXO12
XX BIZYMA
KW Hepati
XX Enzyma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to a novel isolated polynucleotide comprising a nucleotide sequence which is a polymorphic variant of a reference sequence for cytochrome p450, subfamily I (dioxin-inducible), polypeptide I (glaucoma 3, primary infantile), (CPPIBI) gene or its fragment, or a polymorphic variant of a reference sequence for a CYPIBI cDNA or its fragment. The polypeptide of the invention has cytostatic and ophthalmological activity. The polymucleotide may have a use in gene therapy, and antisense gene therapy. The polymorphism and haplotype data of the invention are useful for validating whether CYPIBI is a suitable target for drugs to treat breast cancer and primary congenital glaucoma, screening for such drugs and reducing bias in clinical trials of such drugs. The sequence represents an allele-specific oligonucleotide (ASO) probe, used in the invention to detect polymorphisms in the CYPIBI gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Enzymatic nucleic acid; RNA cleavage; Hepatitis C virus infection; HCV ribozyme; HCV expression; HCV replication; cirrhosis; virucide; liver failure; hepatocellular carcinoma; HCV infection; drug therapy; type I interferon; interferon alpha; interferon beta; cytostatic; type I interferon;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New genetic variants of cytochrome P450, subfamily I dioxin-inducible, polypeptide 1, glaucoma 3, primary infantile gene, CYP1B1 for treatmen and expressing CYP1B1 protein for use in identifying drugs to breast
                                                                                                                                                                                  (ROBE/)
(PAVC/)
                                                                                                                                                                                                                   (BLAT/)
                                                                                                                                                                                                                                                                                                        23-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                               US2002082225-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hepatitis C virus substrate #1037 for HCV hammerhead ribozyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-DEC-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 15 BP; 0 A; 6 C; 1 G; 7 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 15;
                                                                                                                                                                                                                                                                       23-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  interferon gamma; consensus interferon; hepatotropic; antiinflammatory;
substrate; hammerhead ribozyme; HH ribozyme; ss.
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                                                                                                                                                                                MCSWIGGEN J
ROBERTS B.
PAVCO P A.
                                                                                                                                                               MACEJACK D.
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                                                                                                                            Mcswiggen JA,
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86.7%;
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                                                                                                                              Roberts B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВP
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Pred. No. 53;
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                                                                                                                              Pavco PA,
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New ribozymes targeting RNA derived from hepatitis C virus inhibit viral replication and are useful to treat hepatitis C virus infections and cirrhosis, liver failure or hepatocellular carcinoma.

Claim

1;

Page

51;

80pp; English.

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RESULT 25
AAL48035/c
ID AAL480;
XX AAL480;
XX AAL480;
XX AL480;
XX
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                                    The present invention provides the protein, gene and cDNA sequences of human colony stimulating factor 3(granulocyte) CSF3. Also described are single nucleotide polymorphisms (SNPs) identified within these sequences. The sequences can be used in the treatment of neutropenia, promyelocytic leukaemia and haematological disorders. The present sequence is an allele
                                                                                                                                                                                                                                                                                                                                                    New variants of colony stimulating factor 3 (CSF3) isogenes, useful improving efficiency and reliability in the development of drugs for treating diseases associated with CSF3 activity e.g. neutropenia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; colony stimulating factor 3(granulocyte); CSF3; SNP; isogene; chromosome 17q11-12; single nucleotide polymorphism; immunostimulant; neutropenia; promyelocytic leukaemia; haematological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to enzymatic nucleic acids which specifically cleave RNA derived from Hepatitis C virus (HCV). The enzymatic nucleic acid or ribozyme is in a hammerhead (HH) or hairpin (HP) motif where the binding arms comprise sequences complementary to of the substrate sequences defined in the specification. The HCV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-JUN-2000; 2000US-0210380P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kazemi
                                                                                                                                                                                                                                                                           Page 13; 68pp;
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    the coding sequences of
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RESULT 26
RESULT 26
RD ABC12425/c
ID ABC124
XX ABC124
XX Oligon
XX SNP; B
KW Peptid
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Matches 12
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                                                                                                                                                                                                                                                                                     range of diseases including immune system, gastrointestinal, respirator central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC000-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, buwas obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                          Sequence 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Set of oligonucleotides, useful for diagnosis and cell typing,
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                                                                                                                                                                                                                                                               ftp.wipo.int/pub/published_pct_sequences
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Pred.
                                                                                                                            Score 11.4;
Pred. No. 59
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                                                                                                                                                        DB 1; Length 13;
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                                                                                                                                             AAX31546 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oligonucleotide SEQ ID NO 12431 for detecting SNP TSC0002943.
                                                             diagnosis; prognosis; treatment;
                                                                                          Tag sequence; colorectal cancer; pancreatic cancer; colon cancer;
                                                                                                                                                                                                                         21-MAY-1999
                                                                                                                                                                                                                                                                                   AAX31546;
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                                                                                                                                                        of a transcript increased in pancreatic cancer.
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                                                                                                                                                                                                                                                                                            Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBB3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoes; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease;
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                                                                                                                                                  WO200078341-A1
                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                      neovascular condition of the retina;
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                                                                                                                                                                                                                                                                                                                                                                                                                       growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antisense therapy; antiproliferative; anticide antipacity; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pittyriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to a method for ameliorating the effects (skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense
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                                                                                             21-JUN-1999;
                                                                                                                                              21-JUN-2000; 2000WO-AU000693
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                                  (MURD-) MURDOCH
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                                  CHILDRENS RES INST
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5 C; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Edmondson
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Pred. No. 6
                                                                                                                                                                                                                                                                                                                                                                                              retina;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            keloid;
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skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense
                                                             hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood
                                                                                                                                       oligonucleotides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
Sequence 15 BP; 1 A; 6 C; 5 G; 3 T; 0 U; 0 Other;
                                         vessels or any other hyperplasia
                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to a method for ameliorating the effects
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wraight CJ,
                                                                                                                                                                                                                                                                                                                                                                                                                          7; Page
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                                                                                                                                                                                                                                                                                                                                                                                                                        46; 201pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Edmondson
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                                                                                                                                                                                        psoriasis,
                                                                                                                                                                                                                                                                                                              of,
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Query Match Best Local ( Matches 1710 GGCTGCTGACTGA 1722 12; N Similarity GGCTGCTGCCTGA 14 Conservative 23.7%; 0, Score 11.4; Pred. No. 60; Mismatches DВ 1; μ, Length 15; Indels 0; Gaps

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RESULT 31
AAS98675/c
ID AAS986
XX AAS986
XX AAS986
XX COlony
XX Colony
XX Colony
XX Gytost
XX Gy Colony stimulating factor 1 receptor; CSF1R; polymorphic variant; cytostatic; gene therapy; malignant histicytosis; isogene; myeloid malignancy; inflammatory disorder; transgenic animal; haplotype; genotype; human; allele specific oligonucleotide; ASO; probe; ss. Colony stimulating factor 1 receptor (CSF1R) oligonucleotide 26-MAR-2002 AAS98675 standard; DNA; 15 (first entry) #41.

Homo sapiens.

WO200179225-A2

12-APR-2001; 2001WO-US012044.

12-APR-2000; 2000US-0196411P

GENAISSANCE PHARM INC

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2002-075058/10

Novel polymorphic variants of colony stimulating factor 1 receptor useful in studying expression and function of the protein, useful for screening candidate drugs to treat diseases e.g. inflammatory disorders.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CC malignant histiccytosis, mysloid malignancies, and inflammatory disorders CC and the haplotypes can be used to validate CSFIR as a candidate target CC for treating a specific condition or disease predicted to be associated CC with CSFIR activity. Genotyping the CSFIR gene of an individual can also be used in developing diagnostic tests and therapeutic treatments. (I) is CC useful in studying the expression and function of CSFIR, and in CC expressing CSFIR protein for use in screening for candidate drugs to CC treat diseases related to CSFIR activity and in studying the effect of CC the variation on the biological activity of CSFIR as well as on the Dinding affinity of candidate drugs targeting CSFIR as well as on the CC useful in a variety of diagnostic and prognostic formats and therapeutic methods. A transgenic animal is useful in studying expression of the CC ESFIR isogenes in vivo, for in vivo screening and testing of drugs targeted against CSFIR protein, and for testing the efficacy of therapeutic agents and compounds. Allele specific oligonucleotides (ASO) CC earget region. Without requiring any a priori knowledge of the phenotypic effect of any particular CSFIR or happened to the invention provides a CC method for identifying lead compounds that are more likely to show efficacy in clinical trials. This sequence is an allele specific CC described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                         Human; ss; primer; Plasminogen activator; urokinase; PLAU; cancer; cytostatic; serine procease; thrombolytic disorder; isogene; PCR; pulmonary embolism; chromosome 10g24-grear; haplotype; genotype; SNP; single nucleotide polymorphism; thrombolytic; gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             colony stimulating factor 1 receptor (CSF1R) gene, found on The polypeptide are useful for improving the discovery and development of drugs for treating diseases associated with CSF1R activity, e.g.,
                                                                                                                                                                                       14-NOV-2001; 2001WO-US044001
                                                                                                                                                                                                                                                                                                                                                                                                                                             Human PLAU gene, allele specific primer #34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABK96525;
                                                                                                                                                17-NOV-2000; 2000US-0249703P
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                                                                                                                                                                                                                                                                                                         Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 15 BP; 4 A; 4 C; 4 G; 2 T; 0 U; 1 Other;
                                  WPI; 2002-519370/55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 15; Page 15; 164pp; English.
                                                                                                           (GENA-) GENAISSANCE PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1711 GCTGCTGACTGATGT 1725
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    describes a novel isolated polynucleotide (I) h is a polymorphic variant (PV) of a reference ating factor 1 receptor (CSF1R) gene, found on
                                                                       Bentivegna SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1;
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Pred. No. 6
                                                                       Koshy B;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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The invention relates to a polynucleotide comprising a first nucleotide
                                                                                                                                                                               Claim 14; Page 14; 92pp; English.
                                                                                                                                                                                         useful for improving efficiency and reliability in drug development for treating thrombolytic disorders and cancer.
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CC association between a trait and at least one haplotype pair CC of the PLAU gene, an isolated oligonucleotide for detecting a CC polymorphism in the PLAU gene, a recombismant non-human organism CC transformed or transfected with the gene or cDNA, fragments of the CC polynucleotides of at least 10 base pairs encompassing a polymorphic cC site, an isolated polymorphic variant PLAU protein or fragment, an CC isolated monoclonal antibody specific for PLAU, a computer system for CC storing and analysing polymorphism data for the PLAU gene and a genome CC antibology for the PLAU gene. PLAU is useful in screening for drugs CC targeting PLAU that are useful for treating thrombolytic disorders and CC cancers. The methods are useful for treating thrombolytic disorders and CC cancers. The methods are useful for improving the efficiency and CC reliability of the discovery and development of drugs for treating training CC condition of disease associated with PLAU activity. In validating PLAU as a drug condition of disease associated with plau activity. The antibody is CC condition of disease associated with plau activity. The antibody is CC polynucleotides are useful in studying the expression and function of PLAU and in expressing PLAU activity. The gene for PLAU is claused on chromosome loa24-ater. The present sequence is an allale sequence (NS1) comprising a PLAU (plasminogen activator, urokinase, a serine protease) isogene selected from isogenes 1-9 and 11-20 given in the specification, where each isogene comprises the regions of the PLAU gene or cDNA and is further defined by the corresponding sequence of polymorphisms (defining single nucleotide polymorphisms, SNP). Also included are methods of haplotyping/genotyping (and predicting the haplotype/genotype of the PLAU gene of an individual, identifying an located on chromosome 10q24-qter. The present sequence is an allele specific primer used to amplify PLAU polynucleotides with a specific polymorphism the PLAU ä

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Query Match
Best Local Similarity
Watches 12; Conserva
                  1729 GGGAACAGACAGGAG
1 GGGAACAGACGAGRG 15
                                     Conservative
                                            23.7%;
80.0%;
                  1743
                                     1;
                                              Score 11.4;
Pred. No. 60;
                                     Mismatches
                                                      DB 1; Length 15;
                                     2;
                                     Indels
                                     0
                                     Gaps
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Sequence 15

BP; 5

A; 2 C; 7 G; 0 T; 0 U; 1 Other;

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RESULT 33
Human pancreatic cancer SAGE tag #52.
                                  23-APR-2002
                                                                   ABK32500;
                                                                                                    ABK32500 standard; DNA; 15
                                  (first entry)
                                                                                                    ВP
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ABK32500/c
ID ABK325
XX
AC ABK325
XX
DT 23-APR
XX
Human;
KW Human;
KW serial
KW cancer
XX
Homo s
XX
US6333
XX
PD 25-DEC
XX
XX
PR 20-MAY
XX
XX
PR 20-Human; colon cancer; colorectal cancer; pancreatic cancer; SAGE taserial analysis of gene expression; diagnostic; prognostic; probe; cancer marker; ss. 20-MAY-1998; 20-MAY-1998; 25-DEC-2001 US6333152-B1 Homo sapiens 98US-00081646 98US-00081646

(UYJO ) UNIV JOHNS HOPKINS

Genetic variants of Plasminogen activator, Urokinase (PLAU) isogenes,

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC99889, ABF00010-ABF99889, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to an isolated, purified human nucleic acid (I) that has the same sequence as a mRNA found in humans and is a SAGE (serial analysis of gene expression) tag comprising a single stranded probe containing at least 10 consecutive nucleotides. SAGE tags, are diagnostic and prognostic markers of cancer, especially of the colon and pancreas. ABK31900-ABK32770 represent human colon and pancreatic cancer
                                                                                                                                                                              Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and
                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; peptide nucleic acid; cytosine methylation; cardiovascular;
                                                                                                                                        Claim 1; SEQ ID NO 355436; 29pp + Sequence Listing; German.
                                                                                                                                                                   designed to detect methylation status.
                                                                                                                                                                                                                                                  Olek A,
                                                                                                                                                                                                                                                                                                          07-APR-2000; 2000DE-01019173
                                                                                                                                                                                                                                                                                                                                   06-APR-2001; 2001WO-IB000713.
                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                               central nervous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 15
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                                                                                                                                                                                                                         WPI; 2001-657177/75
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12; Conserv
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                                                                                                                                                                                                                                                                              EPIGENOMICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                              system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           <u>,</u>
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Pred. No. 60
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RESULT 35
ABP42717/C
ID ABP427
XX ABP42717/X
AC ABP427
XX 21-PEB
XX Oligon
XX SNP; & 
We peptid
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Best Local S
Matches 11
                                                                                                                                                                                              Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
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                                                                                                                                      Sequence 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; SEQ ID NO 142714; 29pp + Sequence Listing;
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                                                                                                                                                                            ftp.wipo.int/pub/published_pct_sequences
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      1721 GATGTTGAGGG 1731
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                                                                                                                                      BP; 3 A; 7 C; 0 G; 2 T; 0 U; 1 Other;
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Pred. No.
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62;
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59;
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RESULT 36
ABH51281/c
ID ABH512
XX ABH512
XX ABH512
XX ABH512
XX Oligon
XX SNP; as
KW Peptid
KW Centra
XX Homo s
XX HOMO s
XX HOMO s
XX HOMO s
XX WO2001
XX O7-APR
XX 07-APR
XX 07-APR
XX 07-APR
XX 07-APR
XX WPI; 2
APT Olek A
XX WPI; 2
XX WPI; 2
XX WPI; 2
XX WPI; 2
XX Oleim
XX Claim
X
RESULT 37
ABC88045/c
ID ABC880
XX
AC ABC880
XY
T21-FEB
XX
COligon
XX
SNP; 8
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Best Local S
Matches 11
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                                            Oligonucleotide SEQ ID NO 88062 for detecting SNP TSC0022137.
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS;
                                                                                                   21-FEB-2002
                                                                                                                                                     ABC88045;
                                                                                                                                                                                                 ABC88045 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-APR-2000; 2000DE-01019173
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100.0%; Pr
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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07-APR-2000; 2000DE-01019173

06-APR-2001; 2001WO-IB000713

18-OCT-2001 WO200177384-A2 Homo sapiens. SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss;

Oligonucleotide SEQ ID NO 128162 for detecting SNP TSC0032096

21-FEB-2002

(first entry)

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central nervous system; gastrointestinal; respiratory; immune; metabolic

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RESULT 38
ABF281.65/c
ID ABF281
XX ABF281
XX ABF281
XX ABF281
XX ABF281
XX Oligon
XX SNP; 8
XW peptid
XW peptid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                       Claim 1; SEQ ID NO 251257; 29pp + Sequence Listing; German
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This invention describes novel oligonucleotide primers or peptide nucleic

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RESULT 40
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XX SAP; B
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XX SNP; B
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                                                                                          acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC99389, ABF00010-ABE99389, ABH00010-ABH99389 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosin methylation status.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; SEQ ID NO 252383; 29pp + Sequence Listing; German.
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RESULT 41
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RESULT 42
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ID ABF28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
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  ABF28164 standard; DNA; 13 BP
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                                                                                                                                                                          1720 TGATGTTGAGG 1730
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Piepenbrock C,
                                                                                                                                                                                                                                                           Conservative
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100.0%; Pr/
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                                                                                                                                                                                                                                                                                                                                                                       G; S T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                      Score 11;
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Pred. No. 62;
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                                                                                                                                                                                                                                                                                                              Length 13;
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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABI32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; peptide nucleic acid; cytosine methylation; cardiovascular; primer; central nervous system; gastrointestinal; respiratory; immune; metal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                      Oligonucleotide SEQ ID NO 142713 for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 13 BP; 2 A; 0 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; peptide nucleic acid; cytosine methylation; central nervous system; gastrointestinal; re
                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                   21-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABF42716;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABF42716 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     was obtained in electronic format from W
ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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                                                                                                                                                                                                                                                                                                           SNP TSC0035797.
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                                                                                                                                 rimer; CNS;
rimer; 88;
metabolic.
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RESULT 44
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central
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                        Olek A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
WPI; 2001-657177/75
                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; peptide nucleic acid; cytosine methylation; cardiovascular;
                                                                                                      06-APR-2001; 2001WO-IB000713.
                                                                                                                                                         WO200177384-A2
                                                                                                                                                                                                                                                                Oligonucleotide
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                                                                                                                                                                                                                                                                                                                                            ABH52407 standard; DNA; 13
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                                                    (EPIG-)
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                                                                              2000DE-01019173
                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                            system;
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                                                                                                                                                                                                            gastrointestinal;
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The present invention describes nucleic acids (A) that interact stably with a target sequence and contain at least one phosphoro(di)thioate link, having endonuclease activity. (A), and more generally any catalytems.

any catalytic

New nucleic acids that interact, used to treat cancer.

and

optionally cleave, target sequences,

Matulic-Adamic Thompson JD, Reynolds M,

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Beigelman Zwick M, i

1 L, Mcswiggen JA,
Jarvis T, Woolf T,

Karpeisky Haeberli

P, A

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2000-013248/01.

20-APR-1998; 23-JUN-1998; 19-APR-1999;

98US-0082404P. 98US-00103636. 99WO-US008547.

(RIBO-) RIBOZYME PHARM INC.

Claim 79; Page 103; 148pp; English

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RESULT 45
AAA26214/c
ID AAA262
XX AAA262
XX AAA262
XX Oestro
XX Inammer
XX Wo9954
XX W09954
XX Inamer
PF 19-APR
XX 20-APR
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Matches 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010—ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavage; hammerhead ribozyme; hairpin ribozyme; antisense oligonucleotide; gene expression modification; cancer; phosphorothioate; endonuclease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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11; Conserv
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1; Mismatches
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RESULT 46
ABC68262
ID ABC68262
AC ABC68
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CC Because of the high selectivity for targeted RNA, (A) can also be used to CC correlate inhibition of gene expression with alterations in phenotype, CC particularly for identification of therapeutic targets, and as research CC respents (for RNA, in the same way that restriction endomucleases are CC used with DNA). The combination of modifications in (A) improves CC resistance to nucleases, binding affinity and/or activity. AAA23503 to AAA24747 represent oestrogen receptor hammerhead ribozyme sequences, and CC AAA24748 to AAA25992 represent their corresponding target sequences. AAA26105 represent their corresponding target sequences and AAA36107 to AAA26271 represent their corresponding target sequences and AAA36219 to AAA26271 represent other ribozyme sequences and contisense oligonucleotides used in the exemplification of the present
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99989, ABF00010-ABF99989 and ABI00010-ABI82073
                                                                                                                                                                                                                                                                                                                                                                               Set of oligonucleotides, useful for diagnosis and designed to detect single-nucleotide polymorphisms methylation status.
                                                                                                                                                                                                                                                                                                               Claim 1; SEQ ID NO 68279; 29pp +
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; ide nucleic acid; cytosine methylation; cardiovascular; primer; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2001-657177/75
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85.7%;
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1739 AGGAGAAATGC 1749

Matches Query Match Best Local

10;

Conservative

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Similarity

22.1%;

Score 10.6; Pred. No. 70;

DB

<u>.</u> 0

Length 13; Indels

0

Gaps

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Sequence 13

BP; 2 A; 5 C; 0 G; 5 T; 0 U; 1

Other;

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RESULT 47
ABC68263/c
ID ABC682
XX ABC682
XX Oligon
XX SNP; s
KW peptid
XX Centra
XX Homo s
XX Homo s
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XX O1-APR
XX O6-APR
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Best Local S
Matches 10
                                                                                                                  This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF9989, ABF00010-ABF99899, ABF00010-ABF9989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989, ABF0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                            ftp.wipo.int/pub/published_pct_sequences
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Pred. No. 70;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SNP TSC0017813.
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RESULT 48
ABH85901
ID ABH855
XX ABH855
XX ABH855
XX ABH855
XX ABH855
XX ABH855
XX ABH856
XX ABH8
RESULT 49
ABI41640
ID ABI41
XX
AC ABI41
AC ABI41
XX
DT 22-FE
XX
DE Oligo
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Best Local S
Matches 11
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                                   Oligonucleotide primer SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; peptide nucleic acid; cytosine methylation; cardiovascular; central nervous system; gastrointestinal; respiratory; immur
                                                                                                22-FEB-2002 (first entry)
                                                                                                                                                         ABI41640;
                                                                                                                                                                                                                   ABI41640 standard; DNA; 12 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 12 BP; 7 A; 0 C; 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention describes novel oligonucleotide primers or peptide nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; SEQ ID NO 285794; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-APR-2000; 2000DE-01019173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  detect single-nucleotide polymorphisms and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 10.4;
Pred. No. 71
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                                   341613 for detecting SNP TSC0042137.
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primer; ss;
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                                                                                                                                                            Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                            Claim 1;
                                                                                                                                                                                                                                         (EPIG-)
                                                                                                                                                                                                                                                           07-APR-2000; 2000DE-01019173.
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                                                                                                                                                                                                    2001-657177/75
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                                                                                                                                           SEQ ID NO 341613; 29pp + Sequence Listing;
                                                                                                                                                                                                                     Piepenbrock C,
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABF00010-ABF99989989 and ABF00010-ABF99989 and A Sequence 12 BP; 5 A; 0 C; 6 G; 1 T; 0 U; 0 Other; ftp.wipo.int/pub/published\_pct\_sequences

밁 ş Query Match
Best Local Similarity
Matches 11; Conserv 1726 TGAGGGAACAGA 1737 11; 1 TGAGGGAAGAGA 12 Conservative 21.7**%**; 91.7**%**; 0 Score 10.4; Pred. No. 71; Mismatches DB 1; 1; Length 12; Indels 0, Gaps

0;

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RESULT 50
ABI49018/c
ID ABI49018 standard; DNA; 12 BE
XX
AC ABI49018;
XX
DT 22-FEB-2002 (first entry)
XX
DE Oligonucleotide primer SEQ II
XX
SNP; single nucleotide polyma
XX
Extra Suppide nucleic acid; cytosir
XX
OS Suppide nucleic acid; cytosir
XX
OS Suppide nucleic acid; cytosir
XX
Weentral nervous system; gasti
XX
OS Homo sapiens.
XX
PD 18-OCT-2001.
XX
PD 18-OCT-2001.
XX
PF 06-APR-2001; 2000DE-01019173
                                                                                                                                                                                                                                                                        Oligonucleotide primer SEQ ID NO 348991 for detecting
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SNP

TSC0045849

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; 06-APR-2001; 2001WO-IB000713. central nervous system; gastrointestinal;

07-APR-2000; 2000DE-01019173.

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RESULT 51
AB162513/c
ID AB1625
XX AB1625
XX AB1625
XX Oligon
XX SNP; 8
KW SPP; 46
KW Centra
XX Homo 8
XX Homo 9
XX H
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Matches
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                                                                            Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
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                                                                                                                                                                                     WPI; 2001-657177/75
                                                                                                                                                                                                                                      Olek A,
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ide nucleic acid; cytosine methylation; cardiovascular; primer;
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Claim 1; SEQ

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NO 362486;

29pp

+

Sequence

Listing; German

range of diseases including immune system, gastrointestinal, respiratory central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00011-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published\_pct\_sequences

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a

respiratory,

ABC00010

Claim 1;

SEQ ID NO 269262; 29pp + Sequence Listing; German.

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RESULT 52
ABH69285/c
ID ABH692
XX ABH692
XX ABH692
XX Oligon
XX SNP; B
KW peptid
KW Centra
XX Homo s
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                                                                                                                                                                                                                                                                                                                                                                          methylation status.
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ARESULT 53
ABC94621/6
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                                                                                                                                                                                                                                              This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
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RESULT 54 ABC36414

central nervous

aystem;

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; peptide nucleic acid; cytosine methylation; cardiovascular; primer;

gastrointestinal; respiratory; immune;

Oligonucleotide SEQ ID NO 118902 for detecting SNP TSC0029684.

21-FEB-2002

(first

entry)

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ABF18905/c
ID ABF189
XX
AC ABF189
XX
DT 21-FEB
XX
Oligon
XX
KW SNP; s
KW SNP; s
KW peptid
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CS Homo s
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                                                                                         ABF18905
                                                                                                          ABF18905 standard;
                                                                                                                                                                                                                          Sequence 13 BP; 6 A; 0 C; 5 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                             ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                              Claim 1; SEQ ID NO 36431; 29pp + Sequence Listing; German
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Pred. No. 75;
0; Mismatches
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RESULT 56
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                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss;
                                                                                                                                                                                                                                                               Oligonucleotide
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             Olek A,
                                                                                              06-APR-2001; 2001WO-IB000713.
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                                                                  07-APR-2000; 2000DE-01019173
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                                        EPIGENOMICS AG.
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                                                                                                                                                                                                          system; gastrointestinal;
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                                                                                                                                                                                                                                                                                          entry)
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                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; peptide nucleic acid; cytosine methylation; cardiovascu central nervous system; gastrointestinal; respiratory;
                                                                                Claim 1; SEQ ID NO 192691; 29pp + Sequence Listing;
                                                                                                              methylation status.
                                                                                                                          Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosin
                                                                                                                                                                    WPI; 2001-657177/75.
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cardiovascular; primer; ss;
espiratory; immune; metabolic
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This invention describes novel oligonucleotide primers or peptide nucleatid (PNA) oligomers for detecting single nucleotide polymorphisms (S) and cytosine methylation status in chemically pretreated genomic DNA. oligonucleotides are used for diagnosis and/or prognosis of cancer and oligonucleotides are used for diagnosis and/or prognosis of cancer and oligonucleotides are used for diagnosis and/or prognosis of cancer and oligonucleotides are used for diagnosis and/or prognosis of cancer and oligonucleotides are used for diagnosis and/or prognosis of cancer and oligonucleotides are used for diagnosis and/or prognosis of cancer and oligonucleotides are used for diagnosis and/or prognosis of cancer and oligonucleotides are used for diagnosis and/or prognosis of cancer and oligonucleotides are used for diagnosis and/or prognosis of cancer and oligonucleotides are used for diagnosis and/or prognosis of cancer and oligonucleotides are used for diagnosis and/or prognosis of cancer and oligonucleotides are used for diagnosis and/or prognosis of cancer and oligonucleotides are used for diagnosis and/or prognosis of cancer and oligonucleotides are used for diagnosis and/or prognosis of cancer and oligonucleotides are used for diagnosis and/or prognosis of cancer and oligonucleotides are used for diagnosis and/or prognosis of cancer and oligonucleotides are used for diagnosis and/or prognosis of cancer and oligonucleotides are used for diagnosis and/or prognosis of cancer and oligonucleotides are used for diagnosis and/or prognosis of cancer and oligonucleotides are used for diagnosis and/or prognosis of cancer and oligonucleotides are used for diagnosis and oligonucleotides a

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Matches 11;
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RESULT 60
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Best Local Similarity
Matches 11; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                     Sequence 13
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                                                                                                                                                                                                                                                                                                 ftp.wipo.int/pub/published_pct_sequences
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AGTGATGTTGAG 13
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                                                                                                                    Conservative
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                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; car peptide nucleic acid; cytosine methylation; cardiovascular; pri central nervous system; gastrointestinal; respiratory; immune;
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91.7%;
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Pred. No. 75;
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Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                         This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for disgnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence
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                                                                                                                                                                                                                                                                                                                                                                                               Set of oligonucleotides, useful for diagnosis and designed to detect single-nucleotide polymorphisms methylation status.
   represent the oligomers data for this patent did
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Olek A,
                                                                                                                                                                                                                                                                                                                                   Claim 1; SEQ ID NO 78940; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-657177/75
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                                                                                                                                                                                                                                                                                       Sequence 13
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ftp.wipo.int/pub/published_pct_sequences
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TGACGTTGAGGG
                                                                                                                                                                                                                                                                                       BP; 3 A; 6 C; 1 G; 2 T; 0 U; 1 Other;
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                                                                                                                                           Conservative
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91.7%;
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                                                                                                                                                                                 Score 10.4;
Pred. No. 7
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Pred. No. 75;
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RESULT 65
ABF92697/c
ID ABF92697 standard; DNA; 13 BP
XX
AC ABF92697;
XC ABF92697;
XX
DT 22-FEB-2002 (first entry)
XX
D1 22-FEB-2002 (first entry)
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D1 22-FEB-2002 (first entry)
XX
C1 ingonuclectide SEQ ID NO 192
XX
SNP; single nuclectide polymc
KW SNP; single nuclectide polymc
KW SNP; single nuclectide polymc
KW SNP; single nuclectide cytosir
XX
SNP (EPIG-12001; 2001WO-IB000713
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   RESULT 66
ABF56731C
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AC ABF567
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DT 21-FEB
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KW SNP; 8
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Matches 11
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   SNP; single nucleotide polymorphism; human; diagnosis; PNA; peptide nucleic acid; cytosine methylation; cardiovascular;
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                                                                                             Oligonucleotide SEQ ID NO 156728 for detecting SNP TSC0039520
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91.7%;
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Pred. No. 75;
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primer;
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RESULT 67
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                                                                                                                                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                         07-APR-2000; 2000DE-01019173.
                                                                                                                  06-APR-2001; 2001WO-IB000713.
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Pred. No. 75;
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                                                                      Claim 1; SEQ ID NO 118903; 29pp + Sequence Listing; German
                                                                                                                                   Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Olek A,
                                                                                                                                                                                                                                                                                                                                                      (EPIG-)
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                                                                                                                                                                                                                                                                                              Piepenbrock C,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 118903 for detecting SNP TSC0029684.
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Pred. No. 7
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invention describes (PNA) oligomers

for detecting single nucleotide polymorphisms

novel oligonucleotide primers or peptide nucleic detecting single nucleotide polymorphisms (SNP)

Sequence 13

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5 A; 5 C;

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RESULT 69
RESULT 69
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Best Local :
                                                                                                          This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 ABC99989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF9989, ABF00010-ABF99899, ABF00010-ABF9989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF9989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989, ABF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 ABC9989, ABF00010-ABF9989, ABH00010-ABF9989, and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; peptide nucleic acid; cytosine methylation; cardiovascular; primer; central nervous system; gastrointestinal; respiratory; immune; metal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 243408; 29pp + Sequence Listing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-657177/75
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Pred. No. 79
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Best Local
                                                                                                                                                                                                                                   Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
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                                                                                                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; SEQ ID NO 111936; 29pp + Sequence Listing; German
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Olek A,
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ilarity 91.7%;
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Pred. No. 75;
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ABF92696 ID ABF9 XX

ABF92696 standard;

DNA;

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ABH14878
ID ABH1
XX
AC ABH1
XX
ZZ 22-F
DT 22-F
DX Olig
XX
EW SNP,
XW SNP,
XW Cent
XX
EW Cent
XX
FM Cent
XX
FM CO2(

22-FEB-2002

(first entry)

Oligonucleotide SEQ ID NO 214855 for detecting

SNP TSC0052286

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic

WO200177384-A2 Homo sapiens. 문 ঠ

RESULT 72

ABH14878 standard; DNA; 13

BP

RESULT 71

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Matches 11;
                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                      range of diseases including immune system, gastrointestinal, respirate central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00-ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, two sobtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                       Sequence 13 BP; 2 A; 1 C; 6 G; 3 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; SEQ ID NO 192693; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                            ftp.wipo.int/pub/published_pct_sequences
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TGATGTCGAGGG 12
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                                                                                                        0; Mismatches
                                                                                                                                      Score 10.4;
Pred. No. 75;
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RESULT 73
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                                                                           Olek A,
                            WPI; 2001-657177/75
                                                                                                                                                                               07-APR-2000; 2000DE-01019173
                                                                                                                                                                                                                                06-APR-2001; 2001WO-IB000713
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-APR-2000; 2000DE-01019173
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                                                                                                                                                                                                                                                                                                                                                                                                                                       central nervous
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                                                                           Piepenbrock C,
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                                                                                                                                                                                                                                                                                                                                                                                                                                    system;
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                                                                                                                                                                                                                                                                                                                                                                                                                                          immune; metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer;
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Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                           Claim 1; SEQ ID NO 78939; 29pp + Sequence Listing; German
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range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published\_pct\_sequences oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The

Sequence 13 BP; 8 A; 0 C; 4 G; 1 T; 0 U; 0 Other;

Matches Best Query Match Local 11; Similarity Conservative 21.7%; 91.7%; 0; Score 10.4; Pred. No. 7 Mismatches DB 1; Length 13; ۲, Indels 0; Gaps 0

밁 Ş 1736 GACAGGAGAAAT 1747 1 GAAAGGAGAAAT 12

RESULT 74
ABF56730
ID ABF56730 21-FEB-2002 ABF56730, ABF56730 standard; (first DNA; entry) 13 В₽

Oligonucleotide SEQ ID NO 156727 for detecting SNP TSC0039520

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; peptide nucleic acid; cytosine methylation; cardiovascular; primer; central nervous system; gastrointestinal: resniratory. immuno. \_\_\_\_\_ imer; ss; metabolic. CNS;

sapiens.

WO200177384-A2

18-OCT-2001

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS ĀG

P Piepenbrock C, Berlin

2001-657177/75

methylation status. Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and typing, i

Claim 1; SEQ ID NO 156727; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for disgnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The range of diseases including immune system, central nervous system, cardiovascular and oligomers are also used for detecting cell ABC00010

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ABF18907/c
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Best Local Simi
Matches 11;
                Query Match
Best Local (
Best Local Similarity
Matches 11; Conserv
                                                                                                          range of diseases including immune system, gastrointestinal, respirate central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABCO -ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, was obtained in electronic format from WIPO at
                                                                                                                                                                                                                     This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                     Set of oligonuclectides, useful for diagnosis and cell typing, is designed to detect single-nuclectide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                       Claim 1; SEQ ID NO 118904; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-657177/75
                                                                                                                                                                                                                                                                                                                                                                                                                                     Olek A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-FEB-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 13 BP; 6 A; 0 C; 5 G; 1 T; 0 U; 1 Other;
                                                            Sequence 13 BP; 2 A; 6 C; 0 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-APR-2000; 2000DE-01019173.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-APR-2001; 2001WO-IB000713.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ftp.wipo.int/pub/published_pct_sequences
                                                                                           ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (EPIG-)
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Conservative
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             21.7%;
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                  Score 10.4;
Pred. No. 7
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Pred. No. 75;
 Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            detecting SNP TSC0029684.
                  75;
                              DB 1;
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                            Length 13;
Indels
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Gaps
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1721 GATGTTGAGGGA 1732

Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:10810.

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RESULT 77
AAF42671/c
ID AAF42
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AC AAF42
XX
AC AAF42
XX
DT 23-MA
XX
DE Yeast
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ABH58052
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Best Local :
                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                          This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                          23-MAR-2001
                                                    AAF42671;
                                                                            AAF42671 standard;
                                                                                                                                                                                                                                                      Sequence 13 BP; 2 A; 0 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1; SEQ ID NO 258029; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Olek A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-APR-2000; 2000DE-01019173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
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                                                                                                                                                                                                                Local Similarity
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                                                                                                                                                                        1720 TGATGTTGAGGG 1731
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                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       system; gastrointestinal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     acid; cytosine methylation;
                                                                              DNA;
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                                                                                                                                                                                                               21.7%;
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Pred. No. 7
                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                75;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        respiratory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       diagnosis; PNA; cancer; CNS; cardiovascular; primer; ss; espiratory; immune; metabolic.
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RESULT 78
AAD25438
ID AAD25
XX
AC AAD25
XX
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AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention.
                                                                                                                                                                                                                                           Matches
                        AAD25438;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yeast gene coding sequences comprising NORF genes with serial analysis of gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification;
                                                            AAD25438 standard; DNA; 10
                                                                                                                                                                                                                                                                                                                    Sequence 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (UYJO ) UNIV JOHNS HOPKINS.
                                                                                                                                                                                          1719 CTGATGTTGA 1728
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                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                    BP; 4 A;
                                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                           20.8%;
                                                                                                                                                                                                                                                                                                                    3 C; 1
                                                                                                                                                                                                                                                                                                                    G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                           Score 10;
Pred. No.
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                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                           DB 1; Length 10; 72;
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RESULT 79
AAD53533
ID AAD53
XX AAD53
XX
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Best Local S
Matches 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to genetic variants of human gonadotropin-releasing hormone 2 (GNRH2) gene. The invention also relates to compositions and methods for haplotyping and/or genotyping the GNRH2 gene in an individual. Polynucleotides of the invention are useful for studying the expression and function of GNRH2 and in expressing GNRH2 proteins for use in screening candidate drugs to treat diseases related to GNRH2 activity. They are also used in gene therapy. The methods of the invention are useful in determining whether an individual has a haplotype or haplotype pairs. The haplotyping method is useful for improving the efficiency and reliability of several steps in the discovery and development of drugs for treating diseases associated with GNRH2 activity, e.g., reproductive disorders. The present sequence is a primer used for detecting human GNRH2 gene polymorphisms
                                                                                                                                                                                                                                                                       Human; gonadotropin-releasing hormone 2; GNRH2; reproductive disorder; gynaecological; cytostatic; hormonal; target validation; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                             Human GNRH2 gene polymorphism detecting primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAD53533 standard; DNA; 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-MAY-2000; 2000US-0205187P.
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                                                                                                                                     Homo sapiens
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ping; gene therapy; reproductive disorder; polymorphism; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 CTGCTGACTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 1 A; 3 C; 3 G; 3 T; 0 U; 0 Other;
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Conservative
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                                                                                                                                                                                                                                lead compound; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
72;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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WO200294850-A2

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RESULT 80
ABV63381
ID ABV63
XX ABV63
AC ABV63
AC ABV63
AX Human
XX Human
XX Homo
XX Homo
XX Homo
XX PSO'I
XX PS
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Best Local S
Matches 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; skin; dermatological; vulnerary; antipsoriatic; antiseborr immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermapsoriasis; dermatitis; skin cancer; EST; expressed sequence tag;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      expression or increased expression of a particular GNRH2 protein isoform, or an expression vector encoding the isoform may be administered to the patient. Haplotype information is useful in improving the efficiency and output of several steps in a drug discovery and development process, including target validation, identifying lead compounds, and early phase clinical trials. GNRH2 gene is used in gene therapy. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to gonadotropin-releasing hormone 2 (GNRH2) and its nucleic acid sequence. Polymorphic variants of the GNRH2 gene are useful in studying the expression and function of GNRH2, and in expressing GNRH2 proteins for use in screening candidate drugs for treating diseases associated with GNRH2 activity, such as reproductive disorders. Polynucleotides comprising a polymorphic gene variant or fragment may be used for therapeutic purposes, where a patient could benefit from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New gonadotropin-releasing hormone 2 genetic variants having polymorphisms function of, and treating disorders, E
                                                                                                                                                                                                                                                                                                                                                WO200253774-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human skin
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     WPI; 2002-590638/63
                                                           Petersohn D,
                                                                                                                                                                        03-JAN-2001; 2001DE-01000127
                                                                                                                                                                                                                                20-DEC-2001; 2001WO-EP015179
                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-OCT-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABV63381 standard; cDNA; 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1712 CTGCTGACTG 1721
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        primer used for detecting human GNRH2 gene polymorphisms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10;
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                                                                                                                   HENKEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kliem SE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EST 1167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 1
                                                           Conradt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Α.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nandabalan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3 C; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20.8%;
                                                           ζ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 g hormone 2 (GNRH2) polypeptide encoded by olymorphisms in the GNRH2 gene, for studying disorders, such as, reproductive disorders.
                                                              Hofmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ۲
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1; Length 10, 72;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antiseborrhaeic;
neurodermatitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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Query Match
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Matches 10
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                                                                                                                                 disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; attopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma, and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (EST) of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                e.g. skin cancer.
                                                                                       Sequence 11 BP; 2 A; 2 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 57; 1345pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     In vitro identification of skin-expressed homeostasis and identifying cosmetic or pl
                       Similarity
                                            20.8%;
                       100.0%;
                       Score 10;
Pred. No.
                       DB 1;
76;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ed genes, useful for determini pharmaceutical agents against
                                            Length 11,
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RESULT 81
ABV70802
ID ABV70802
ID ABV70802
ID ABV7
XX ABV7
XX Huma
XX Huma
XX Homm
XX D8001
OX T10
OX D8001
OX T10
OX T1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
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The invention relates to in vitro identification (M1) of genes expresse in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or

e.g.

cancer.

In vitro identification of a homeostasis and identifying

skin-expressed genes, useful for g cosmetic or pharmaceutical agent

agents

against

expressed

determining

2002-590638/63

Petersohn

'n

Conradt KGAA

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Hofmann

(HENK ) HENKEL

03-JAN-2001; 2001DE-01000127 20-DEC-2001; 2001WO-EP015179 11-JUL-2002

Claim 24;

Page 275; 1345pp;

German.

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RESULT 82
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Matches
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                                                                                The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; inhthyosis; atopic dermatitis, acne, sebornhea; hupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (EST) of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        disorders, specifically neurodermatitis; sunburn; psoriasis; sclerodermatichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma, basal cell carcinoma; and carcinoma or sarcoma of the sequence is that of a human expressed sequence tag
                                                                                                                                                                                                                                                                                                                   In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 11 BP;
                                                                                                                                                                                                                                                                         Disclosure; Page 182; 1345pp; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    promotes skin homeostasis or that can be used for treating skin
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                Local
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  Similarity
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  Conservative
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100.0%; Pr
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100.0%; 100
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                20.8%; Score 10; 100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4 G;
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                                                           <u>ი</u>
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Pred. No.
                                                          6 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3 T;
    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                            <u>۲</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0 U;
                                                          0 U; 0 Other;
                DB 1; Length 11; 76;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       76;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1;
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    0
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    Gaps
    0
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RESULT 84
ABI79664
ID ABI7
XX
AC ABI7
XX
AC ABI7
XX
XX
XX
XX
XX
XX

ABI79664 standard;

DNA; 12

ВP

0

22-FEB-2002

(first entry)

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RESULT 83
ABV64816/c
ID ABV648
XX Human;
XX Human;
XX Homo s
XX Homo s
XX Homo s
XX WO2002
XX IN JUL
XX IN VIT
PT Peters
XX IN VIT
PT Homeos
PT e.g. [
XX PAI HENK
XX PAI HENC
XX IN VIT
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XX PAI HENC
XX ABV61;
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XX PAI HENC
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밁
                                                         S
                                                                                                                              Matches
                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression.

(M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                                                                                                                                            Sequence 11 BP; 5 A; 1 C; 3
                                                                                                                                                                                                                                                                                                                                                 disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the present sequence is that of a human expressed sequence tag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 97; 1345pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-DEC-2001; 2001WO-EP015179
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           In vitro identification of skin-expressed genes, useful for determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (HENK ) HENKEL KGAA.
                                                                                                                                                            Local
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11
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                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                             invention
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                                                                                                                                                                                                                                                            G; 2 T; 0 U; 0 Other;
                                                                                                                                                            Score 10;
Pred. No.
                                                                                                                                 Mismatches
                                                                                                                                                            DB 1; Length 11; 76;
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RESULT 85
ABH71192
ID ABH71
XX ABH71
XX
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Best Local S
Matches 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC99989, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; peptide nucleic acid; cytosine methylation; cardiovascular; central nervous system; gastrointestinal; respiratory; immur
                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss;
                                                                                                                                                                                                                                                                                                                                                                Oligonucleotide
06-APR-2001; 2001WO-IB000713.
                                                                                                                                                                                                                                                                                                                                                                                                                            22-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABH71192;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 12 BP; 6 A; 0 C; 4 G; 2 T; 0 U; 0 Other;
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                                                               18-OCT-2001.
                                                                                                                                                                                   Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1739 AGGAGAAATG 1748
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Piepenbrock C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                         system; gastrointestinal;
                                                                                                                                                                                                                                                                                                                                                                primer SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 system; gastrointestinal; respiratory; immune; metabolic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
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80;
                                                                                                                                                                                                                                         respiratory;
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                                                                                                                                                                                                                                            immune; metabolic
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                             Claim 1; SEQ ID NO 271169; 29pp + Sequence Listing; German
                                                                                                                                                                                                                                                                                                                                                   designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                            Set of oligonucleotides, useful for diagnosis and cell typing,
                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-657177/75
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Matches Query Match Sequence 12 BP; 2 A; 0 C; 5 G; 5 T; 0 U; 0 Other; Local 1722 ATGTTGAGGG 1731 10; Similarity Conservative 20.8%; 100.0%; 0 Score 10; Pred. No. Mismatches ,08 DB 1; Length 12; 0 Indels 0 Gaps 0,

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RESULT 86
ABH80356/c
ID ABH803
XX ABH803
XX ABH803
XX 22-FEB
XX Oligon
XX SNP; s
KW peptid
KW centra
XX Homo s
XX Homo s
XX HOTO S
YN WO2001
XX WO2001
XX UFF O6-APR
XX (EPIG-
XX (EPIG-
XX Olek A
XX WPI; 2
XX Set of
PT Set of
PT design
PT methyl
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                                                                                                                                                                                                                  Oligonucleotide primer SEQ ID
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                                                                                                                                                                                                                                                                                                                  ATGTTGAGGG
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                                                                                                                                                                                                                   NO 280359 for detecting
                                                                                                                                                                                                                    SNP TSC0008516.
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SNP; single nucleotide polymorphism; human; diagnosis; peptide nucleic acid; cytosine methylation; cardiovascu central nervous system; gastrointestinal; respiratory; diagnosis; PNA; cancer; CNS; cardiovascular; primer; ss; metabolic.

Homo sapiens.

WO200177384-A2

18-OCT-2001

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS ÃG.

Piepenbrock C, Berlin ×

Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1;

SEQ

ij ŏ

280359; 29pp + Sequence Listing; German.

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RESULT 87
ABH87503
ID ABH87503
AC ABH87
XX ABH877
XX ABH87
XX ABH87
XX Olige
XX SNP;
XW Pepti
XW Centt
XX Homo
XX O7-Ai
XX O7-Ai
XX O1-Ai
XX O1-
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Best Local :
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosts and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABC9989, ABF00010-ABC998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; SEQ ID NO 287496; 29pp + Sequence Listing; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    methylation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; SEQ ID NO 301335; 29pp + Sequence Listing; German.
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1739 AGGAGAAATG
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                                                                                                                                               BP; S A; 0 C; 4 G;
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100.0%; F1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA, attached to a solid support, and reporter DNA, where either a part of the anchor DNA or reporter DNA is designed to form a triple-strand structure with part of the test sequence. Triples formation results in displacement of the reporter DNA which is detected as an indication of the presence of the DNA test sequence. The method is used to detect DNA sequences, particularly for identification of bacteria (by detecting genes for ribosomal RNA) in clinical samples, but also detection of oncogenes and Hepatitis B virus. (Updated on 27-AUG-2003 to correct OS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence represents a polynucleotide that is able to form a triple helix with a double stranded sequence. Cytosine bases in the present can be replaced with 5-methylcytosine for increased triplex stability. The present sequence is used in the assay of the invention, where it can be part of the anchor DNA or reporter DNA sequence. The assay comprises adding a sample containing double-stranded DNA test sequences to an aqueous medium containing at least one complex of anchor DNA attractions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Assay of genetic sequences based on triplex formation from double stranded analyte - and hybrid of anchor and reporter sequences, with reporter released if triplex formation occurs, used e.g. to identify
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24-MAR-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX14857;
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                                                                                                                          ABF28166 standard; DNA; 13
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                                                                                                                                                                                                                                                                                                                                        1727 GAGGGAACAG 1736
                                                                                                                                                                                                                                                                                                                                                                                                        10;
                                                                                                                                                                                                                                                                                 12 GAGGGAACAG 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PROFILE DIAGNOSTIC SCI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 0 A; 7 C; 1 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Col 21-22; 168pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Wang
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RESULT 91
ABH24633/c
ID ABH246
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oligonucleotide SEQ ID NO 128163 for detecting SNP TSC0032096
                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; peptide nucleic acid; cytosine methylation; cardiovascular; primer; central nervous system; gastrointestinal; respiratory; immune; metal
                                                                                                                                                                                            Oligonucleotide SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                       WO200177384-A2
                                                                                      Homo sapiens
                                                                                                                                                                                                                              22-FEB-2002
                                                                                                                                                                                                                                                                ABH24633
                                                                                                                                                                                                                                                                                               ABH24633 standard; DNA; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID NO 128163; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Set of oligonucleotides, useful for diagnosis and cell typing,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-657177/75
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                                                                                                                                                                                         ID NO 224610 for detecting SNP TSC0054745.
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Pred. No.
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Matches 10; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
designed to detect methylation status.
                        Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      data for this patent did not form part of the printed specification, was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                 07-APR-2000; 2000DE-01019173
                                                                                                                                                                                                                                                                                                                  06-APR-2001; 2001WO-IB000713.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligonucleotide SEQ ID NO 187822 for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-FEB-2002
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Pred. No.
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84;
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range of diseases including immune system, gastrointestinal, respirator central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC000-ABC9989, ABF00010-ABF9989, ABH00010-ABH9988 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, bu was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                              acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNI and cytosine methylation status in chemically pretreated genomic DNA. oligonucleotides are used for diagnosis and/or prognosis of cancer and
                                                                                                                                                                                                                                                                                                                                                                               Claim 1; SEQ ID NO 187822; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                invention
BP; 4 A; 7 C; 0 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                           describes novel oligonucleotide primers or peptide nucleic
                                                                                                                                                                                                                               respiratory,
                                                                                                                                                                               ABC00010
                                                                                                                                                                                                                                                                                                        (SNP)
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RESULT
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Best Local (
                                                                                                                   Matches
21-FEB-2002
                  ABC95357;
                                   ABC95357 standard; DNA; 13
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                                                                                11 TGTTGAGGGA
                                                                                                                  10;
                                                                                                                           Similarity
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(first
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entry)
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                                                                                                                   Score 10; DB; Pred. No. 84; 0; Mismatches
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                                                                                                                                   DB 1;
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뭐 S

Sequence 13

SNP; single nucleotide polymorphism; human; peptide nucleic acid; cytosine methylation; central nervous system; gastrointestinal; re n; cardiovascular; primer; ss;
respiratory; immune; metabolic. diagnosis; PNA; cancer; CNS;

Oligonucleotide SEQ ID NO 95374

for detecting SNP TSC0023742.

Homo sapiens.

WO200177384-A2

18-OCT-2001.

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS AG.

Olek A, Piepenbrock C, Berlin 7

2001-657177/75.

Set of oligonucleotides, designed to detect single methylation status. onucleotides, useful for diagnosis and detect single-nucleotide polymorphisms cell typing, s and cytosine

Claim 1; SEQ ID NO 95374; 29pp + Sequence Listing; German

ABC95357

ID ABC9

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AC ABC9

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AC ABC9

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                                                                                                                                            This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immediate the control of the 
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                                                                                                                      Oligonucleotide SEQ ID NO 206483 for detecting SNP TSC0050541.
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; peptide nucleic acid; cytosine methylation; cardiovascular; primer;

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                                                                                                                                                                                                                                                This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                      Sequence 13
                                                                                                  ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                        Claim 1; SEQ ID NO 197339; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                            methylation
                                                                                                                                                                                                                                                                                                                                                                                            Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine
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10;
                  Similarity
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                                                                      BP; 3 A; 0 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                             status
Conservative
                20.8%;
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<u>,</u>
                  Score 10;
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   Mismatches
                  DB 1;
84;
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                                 Length 13
   Indels
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   Gaps
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RESULT 98
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ID ABH545
XX ABH545
XX ABH545
XX Oligon
XX SNP; s
KW peptid
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XX Homo s
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CC This i Oligonucleotide SEQ ID NO 254570 for detecting SNP TSC0062063 22-FEB-2002 ABH54593; ABH54593 standard; (first DNA; entry) 13 ВР

central nervous system; gastrointestinal; respiratory; immune; SNP; single nucleotide polymorphism; human; peptide nucleic acid; cytosine methylation; diagnosis; PNA; cancer;
cardiovascular; primer; 88;

Homo sapiens.

18-OCT-2001

06-APR-2001; 2001WO-IB000713

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS Ą

Olek A, Piepenbrock C, Berlin ζ.

WPI; 2001-657177/75

Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosin methylation status. cytosine 18

Claim 1; SEQ ID NO 254570; invention describes (PNA) oligomers for novel oligonucleotide primers or peptide nucleic detecting single nucleotide polymorphisms (SNP) 29pp + Sequence Listing; German.

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RESULT 99
ABP97343/c
ID ABP9733
XX ABP973
XX ABP973
XX BP973
XX BP973
XX BP973
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XX BP973
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XX WPI; 2
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                                                                                  and cytosinė methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABC9989, ABF00010-ABC9989, and MBI00010-ABC9989 are present the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1; SEQ ID NO 197340; 29pp + Sequence Listing; German.
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                                                             ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 197340 for detecting SNP TSC0048566.
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                                                                                                                                                                                                                                                                                             This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
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RESULT 102
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                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
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                                                                                                                                                                                                                                                      Oligonucleotide SEQ ID NO 224609 for detecting SNP TSC0054745.
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         WO200177384-A2
                                                                    Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                    ABH24632;
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RESULT 103
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Matches 10; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; peptide nucleic acid; cytosine methylation; cardiovascular; primer; central nervous system; gastrointestinal; respiratory; immune; metal
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                                             Olek A, Piepenbrock C,
                                                                                                        07-APR-2000; 2000DE-01019173
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RESULT 104
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                                                                            The present invention describes a method for the replication of a DNA in which only a specific DNA contained in a mixture of DNAs is replicated by using a polymerase chain reaction (PCR), comprising a primer acting as a replication point to the DNAs present in mixture and a sequence-specific substance which adheres only to the DNA. The method can be used for giving important informations in the mechanism of cancerisation of cells in which many similar genes interact complicatedly. The present sequence
                                                                                                                                                                                                                                                                                                                                                                      Replication method of DNA and apparatus for designing polymerase reactions for giving important information in the mechanism of cancerization of cells.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-JAN-1999;
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                                                        represents a primer which is used in the exemplification of
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                                                                                                                                                                                                                                                                                                           Page 5; 13pp; Japanese.
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                                                                                                                                                                                                                   Sequence 13
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                                                                                                                                                                                                                   BP; 2 A; 7 C; 0 G; 4 T; 0 U; 0 Other;
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Pred. No. 8
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                                                                                                                                                          DB 1; Length 13;
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cardiovascular;
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RESULT 106 ABH34097/c

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RESULT 107
ABC20825/c
ID ABC208
XX ABC208
AC ABC208
XX Oligon
XX Oligon
XX SNP; s
KW SNP; s
KW SNP; s
KW peptid
KW centra
XX OS Homo s
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                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; peptide nucleic acid; cytosine methylation; cardiovascular; central nervous system; gastrointestinal; respiratory; immun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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    Homo sapiens
                                                                                                 Oligonucleotide
                                                                                                                                20-FEB-2002
                                                                                                                                                               ABC20825
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                                                                                            SEQ ID NO 20842 for detecting SNP TSC0004233.
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                                    respiratory; immune;
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Olek A,

Piepenbrock C,

Berlin K;

07-APR-2000; 2000DE-01019173 06-APR-2001; 2001WO-IB000713

(EPIG-) EPIGENOMICS

18-OCT-2001 WO200177384-A2

metabolic.

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RESULT 108
ABC07406
ID ABC074
XX ABC074
AC ABC074
XX Oligon
XX SNP; sq
KW SNP; d
KW Peptid
KW centra
XX Centra
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; peptide nucleic acid; cytosine methylation; cardiovascular; primer; central nervous system; gastrointestinal; respiratory; immune; metal
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Pred. No. 89;
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RESULT 109
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XX SNP; as yeptid
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OS Homo a
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This invention describes novel oligonucleotide primers or peptide nuc acid (PNA) oligomers for detecting single nucleotide polymorphisms (S; and cytosine methylation status in chemically pretreated genomic DNA. oligonucleotides are used for diagnosis and/or prognosis of cancer an range of diseases including immune system, gastrointestinal, respirat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                    Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; peptide nucleic acid; cytosine methylation; cardiovascular;
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                                                                                                            Claim 1; SEQ ID NO 107045; 29pp + Sequence Listing; German.
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RESULT 110
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XX SNP; s
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                                                                                                                                                                                                                                                                                                           This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; SEQ ID NO 87951; 29pp + Sequence Listing; German.
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Query Match 20.. Best Local Similarity 84. Matches 11; Conservative

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RESULT 111
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ID ABF216
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XX Oligon
XX SNP; g
KW Peptid
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Matches 11
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22-FEB-2002
                                                         ABH20206
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RESULT 113
ABH34096
ID ABH340
XX ABH340
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XX OBH340
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XX OBH340
XX OBH340
XX SNP; S WW Peptid
XW Peptid
XW Centra
XX Homo S
XX Hom

22-FEB-2002 ABH34096;

(first

entry)

ABH34096 standard;

DNA;

18-OCT-2001 WO200177384-A2 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Oligonucleotide SEQ ID NO 234073 for detecting SNP TSC0057120.

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1718 ACTGATGTTGAGG 1730

ATTGATGTTTAGG 13

Query Match
Best Local S
Matches 11

Similarity

20.4%;

Score 9.8; Pred. No. (

89;

DB 1; Length 13; <u>ب</u>

Conservative

<u>.</u>

Mismatches

Indels

0

Gaps

0;

Sequence 13

BP; 3

A; 0 C; 4 G;

6 T; 0 U; 0 Other;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                     Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                                                                                  Claim 1; SEQ ID NO 220183; 29pp + Sequence Listing; German.
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RESULT 114
ABH42048
ID ABH420
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XX SNP;
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                                      Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine
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range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a

Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.

Claim 1; SEQ ID NO 48779; 29pp + Sequence Listing;

German

typing, cytosine

Olek

Piepenbrock C,

Berlin

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2001-657177/75

(EPIG-)

EPIGENOMICS

07-APR-2000; 2000DE-01019173 06-APR-2001; 2001WO-IB000713. 18-OCT-2001 WO200177384-A2 Homo sapiens.

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RESULT 115
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                                                                                                                                                                                                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
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                                                                                                                                                                                                                                                                                            This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosts and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9989, ABF00010-ABC9989, ABF00010-ABC9989, ABF00010-ABC9989, ABF00010-ABC9989, ABF00010-BC273 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                 Sequence 13 BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; SEQ ID NO 12434; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-APR-2000; 2000DE-01019173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            was obtained in electronic format from WIPO at
ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     single nucleotide polymorphism; human; diagnosis; PNA; ide nucleic acid; cytosine methylation; cardiovascular;
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SNP; single nucleotide polymorphism; human; peptide nucleic acid; cytosine methylation;
                                                                                       Oligonucleotide SEQ ID NO 41644 for detecting SNP TSC0012495.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABF00010-ABF9989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
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                                                                                                                                                                                                                                      Oligonucleotide SEQ ID NO 224221 for detecting SNP TSC0054636.
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                                                                                                                         Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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Claim 1; SEQ

invention describes novel oligonucleotide primers or peptide nucleic (PNA) oligomers for detecting single nucleotide polymorphisms (SNP)

ID NO 186803; 29pp + Sequence Listing; German

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RESULT 121
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                                                              This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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RESULT 123 ABH42049/c ID ABH420 XX

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                                       Sequence 13
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ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-APR-2000; 2000DE-01019173
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                                                      1744 AAATGCATCCATT 1756
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AAATACATACATT 13
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                                                                                                                  Conservative
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Best Local 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                         Oligonucleotide
                                                                                                                                                                                                                                                          Sequence 13 BP; 3 A; 4 C; 0 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                          methylation status.
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                Homo sapiens
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                                                                                                                                                                              TTTAGGGAAAAGA 1
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                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                 onucleotides, useful for diagnosis and cell typing, is detect single-nucleotide polymorphisms and cytosine
                                                                                          (first entry)
                                                                     SEQ ID NO 250249 for detecting SNP TSC0061098.
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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                      Olek A,
                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                                                                                                                                                                                              Oligonucleotide SEQ ID NO 74891 for detecting SNP TSC0019229.
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                                                                                                                                                                                                             Homo sapiens.
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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010
                                                                                                                                                                                                                                                                                               designed to methylation
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                                                                                                                                                                            invention describes novel oligonucleotide primers or peptide nucleic
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RESULT 127
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Best Local
                                                                     This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99089, ABH00010-ABF99089, ABH00010-ABF9908999, ABH00010-ABF99089, ABH00010-ABF99089, ABH00010-ABF99089, ABH00010-ABF99089, ABH00010-ABF99089, ABH00010-ABF99089, ABH00010-ABF99089, ABH00010-ABF99089, ABH00010-ABF99089, ABH00010-ABF990899, ABH00010-ABF99089, ABH00010-ABF99089, ABH00010-ABF99089, ABH
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Sequence 13
                                                                                                                                                                                                                                                                                                                                                                                           Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                              methylation
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BP; 6 A; 4 C; 0 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                              status
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Pred. No. 89;
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RESULT 128
ABH10408
ID ABH104
XX ABH104
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XX BH104
XX BNP; B
DE Oligon
XX SNP; B
KW Peptid
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   RESULT 129
ABF86807/c
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Matches 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a reange of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation, ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
Oligonucleotide SEQ ID NO 186804 for detecting SNP TSC0046049.
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                                                                                                                                 ABF86807;
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RESULT 130
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                                                                                                                                                                                                                                                                                                                                                                                                   Oligonucleotide SEQ ID NO 188828 for detecting SNP TSC0046484.
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                                                                                                                                                                                                                                                                               central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                        Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
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                                                                                                                                                                                                                                            Olek A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oligonucleotide SEQ ID NO 188830 for detecting SNP TSC0046484
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Claim 1; SEQ ID NO 188830; 29pp + Sequence Listing; German

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9988, ABF00010-ABF9988, ABH00010-ABH9988 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                         This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for disgnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                          Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                   Sequence 13 BP;
                                                                                                                                                                                                                                                                                                   Claim 1; SEQ ID NO 20841; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                             methylation status.
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nes 11; Conserv
                       1739 AGGAGAAATGCAT 1751
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RESULT 134

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ABC97855;
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DT
21-FEB-2002 (first of ABC97855;
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SNP; single nucleotic scid of Central nervous system of ABC9784-A2.
XX
DR
DB
18-OCT-2001.
XX
OS Homo sapiens.
XX
PN
W0200177384-A2.
XX
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06-APR-2001; 2001WO-XX
PR
07-APR-2001; 2000DE-XX
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07-APR-2001; 2000DE-XX
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07-APR-2001; 2000DE-XX
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01ek A, Piepenbrock
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WP1; 2001-657177/75.
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Cestion of diseases in methylation status.
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Claim 1; SEQ ID NO 9
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acid (PNA) oligomers
and cytosine methyla
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KW peptid
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Matches 11
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                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss;
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                                                central nervous system;
                                                                                                                                                                                                                Oligonucleotide SEQ ID NO 48780
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RESULT 136
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  Olek A,
                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss;
                                                                                                              07-APR-2000; 2000DE-01019173.
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                                                                                                                                                                   2001WO-IB000713.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                 system; gastrointestinal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA;
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     Berlin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 9.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            for detecting SNP TSC0002151.
  ζ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                    respiratory;
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d cytosine
                                                                                                                                                                                                                                                                                                                                                                                       immune; metabolic
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                                                                     acid (PNA) oligomers for detecting single nucleotide primers or peptide nucleic and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosts and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABE99989, ABF00010-ABF99899, ABF00010-ABF99899 and ABI00010-ABF39899 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                        Sequence 13 BP; 2 A; 7 C; 0 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 7398;
                                                                                                                                                                                                                                                                                                                                                                                            methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                            Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosin
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20.4%;
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  Score 9.8;
Pred. No. 8
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                    Length 13
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Matches 11; 1720 TGATGTTGAGGGA 1732 13 Similarity Conservative 0; Mismatches 2; Indels <u>,</u> Gaps

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RESULT 137
ABF183
ID ABF183
XX ABF183
AC ABF18 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Oligonucleotide SEQ ID NO 118395 for detecting 21-FEB-2002 ABF18398 standard; DNA; 13 (first entry) BP. SNP TSC0029588.

Homo sapiens.

WO200177384-A2

18-OCT-2001

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS Ã

Piepenbrock C, Berlin 7

WPI; 2001-657177/75.

Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosin methylation status. designed to methylation cytosine

Claim 1; SEQ ID NO 118395; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a

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RESULT 138
ABF21699/c
ID ABF216
XX ABF216
XX ABF216
XX ABF216
XX Y PEB
XX Oligon
XX SNP; s
XW Peptid
XX Centra
XX Homo s
XX WO2001
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XX O1ek A
XX WFI; 2
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PT design
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                                                                                                                                        This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABC99989, ABF00010-ABC99989, aBH00010-ABC99989, and ABI00010-ABC998973 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Olek A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
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                                                                          Sequence 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Piepenbrock C,
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                                                                        A; 6 C; 0
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20.4%;
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                                                                        G; 3 T; 0 U; 0 Other;
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Score 9.8; D
Pred. No. 89;
                         DB 1; Length 13;
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RESULT 139
ABP46516
ID ABP465
XX ABP465
XX ABP465
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XX BP7
AC Oligon
XX SNP; s
KW Peptid
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Matches 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989, ABH00010-ABH99989, ABH0010-ABH99989, ABH0010-ABH999989, ABH0010-ABH999989, ABH0010-ABH999989, ABH0010-ABH999989, ABH0010-ABH999989, ABH0010-ABH999989, ABH0010-ABH999989, ABH0
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                                                                                                                                                                                                                                                                                                                             Sequence 13 BP; 6 A; 0 C; 4 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                 84.6%;
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Pred. No. 89;
                                                                                                                                                              Mismatches
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                                                                                                                                                              Indels
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RESULT 140
ABF51962
ID ABF519
XX
AC ABF519
XX

ABF51962 standard;

DNA;

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ABF51962

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RESULT 141
ABF51963/c
ID ABF519
XX
ABF519
XX
AC ABF519
DT 21-FEE
XX
Oligon
XX
SNP; E
KW SNP; E
KW peptic
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XX
OS Homo E
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XX
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Matches 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory,
    18-OCT-2001
                                   WO200177384-A2
                                                                                            peptide nucleic central nervous
                                                                                                                                                         Oligonucleotide
                                                                                                                                                                                                                      ABF51963
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 13 BP; 4 A; 0 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  designed to detect methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                Homo sapiens.
                                                                                                                                                                                        21-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                   Local Sim-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; SEQ ID NO 151959; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Set of oligonucleotides,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-657177/75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Olek A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-APR-2000; 2000DE-01019173.
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                                                                                           single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; ide nucleic acid; cytosine methylation; cardiovascular; primer; ss; ral nervous system; gastrointestinal; respiratory; immune; metabolic
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                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nnucleotides, useful for diagnosis and cell typing, detect single-nucleotide polymorphisms and cytosin
                                                                                                                                                                                        (first entry)
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                                                                                                                                                         SEQ ID NO 151960
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                                                                                                                                                                                                                                                                                                                                                                                                           Pred.
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                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                        for detecting
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                                                                                                                                                                                                                                                                                                                                                                                                           89;
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                                                                                                                                                         SNP TSC0038398.
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                                                                                                                                                                                                                                                                                                                                                                                                                      Length 13;
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                                                                                            primer; ss;
ne; metabolic
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Matches 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI;
                                 ABF02354 standard;
                                                                                                                                                                                                                                                Sequence 13 BP; 5 A; 4 C; 0 G; 4 T; 0 U; 0 Other;
ABF02354;
                                                                                                                                                                                                                                                                                 was obtained in electronic format from W
ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID NO 151960; 29pp + Sequence Listing; German.
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                                                                                                                                                1720 TGATGTTGAGGGA 1732
                                                                                                                                                                                                                                                                                                                                                                                                                                                      invention describes novel oligonucleotide primers or peptide nucleic (PNA) oligomers for detecting single nucleotide polymorphisms (SNP)
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                                                                                                                  13
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                                                                                                                  TAATTTTGAGGGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Piepenbrock C,
                                                                                                                                                                                 Conservative
                                 DNA; 13
                                                                                                                                                                                                 20.4%;
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                                 ВÞ
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Pred. No. 8
                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                    , 68
                                                                                                                                                                                                                  DB
                                                                                                                                                                                                                  1; Length 13;
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cytosine
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ABF02354
II ABF0
XX ABF(
XX ABF(
XX C)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oligonucleotide SEQ ID NO 102351 for detecting
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-OCT-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        central nervous system;
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                                                                                                                                                                                                                                                                                                                                   (EPIG-)
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ytosine methylation; cardiovascular; primer; ss;
gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                           Berlin
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Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and

typing, is cytosine

WPI; 2001-657177/75.

methylation status

1; SEQ

ID NO 102351; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomers are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published\_pct\_sequences

Matches 11; Query Match

Local

Similarity

20.4%;

Score 9.8; Pred. No. 8 Mismatches , 68 멂 1;

Conservative

<u>.</u>

2

0

Gaps

0

Length 13; Indels

Sequence 13 'BP; 5 A; 0 C; 4 G; 4 T; 0 U; 0 Other;

0

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RESULT 143
ABC41626
ID ABC416
AC ABC416
AX SNP;
COLIGORY
AND ABC416
AX SNP;
KW Pepti
KW Centr
AX Centr
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    This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for disgnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  designed to methylation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                      ID NO 41643;
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1724

GTTGAGGGAACAG 1736

Similarity

Conservative

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Indels

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Gaps

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RESULT 144
ABC97854
ID ABC978
XX ABC978
XX ABC978
XX 21-FEB
XX SNP; see
XX SNP
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Query Match
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Matches 11
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                                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-APR-2000; 2000DE-01019173
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                                                                                                             BP; 3
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Score 9.8; DB Pred. No. 89; 0; Mismatches
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Pred. No. 89;
                                                                                                             2 T; 0 U;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diagnosis and cell typing, a polymorphisms and cytosine
                                                           DΒ
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                                                                                                                   0 Other;
                                                        1; Length 13;
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SNP;
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ABF14030/c
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 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer;
                           Oligonucleotide SEQ ID NO 188827 for detecting SNP TSC0046484.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and
                                                        22-FEB-2002
                                                                                   ABF88830;
                                                                                                             ABF88830 standard; DNA; 13 BP
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                                                                                                                                                                                                                                                                                              0 C; 4 G;
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                                                                                                                                                                                                                                                                 Score 9.8;
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                                                                                                                                                                                                                                                                                               0 Other;
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RESULT 147
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Best Local &
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                               07-APR-2000; 2000DE-01019173
                                                                                        06-APR-2001; 2001WO-IB000713.
                                                                                                                                                                                                                                                                                                                                  central nervous system; gastrointestinal; respiratory; immune;
                                                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; peptide nucleic acid; cytosine methylation; cardiovascular; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oligonuclectide SEQ ID NO 244511 for detecting SNP TSC0059697
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central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
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Pred. No. 89;
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RESULT 148
ABC53910
ID ABC539
XX ABC539
XX 21-FEB
XX 2NP; g
KW SNP; g
KW peptid
KW centra
XX WO2001
XX WO2001
XX WO2001
XX WO2001
XX GEPIG-
PI 06-APR
XX (EPIG-
PX 07-APR
XX (EPIG-
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Best Local S
Matches 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIFO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                            Set of oligonucleotides, useful for designed to detect single-nucleotide methylation status.
                                                                                                                                                                                                                                                                            Olek A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
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                                                                                                                                                                                                                        WPI; 2001-657177/75
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oligonucleotide SEQ ID NO 53927 for detecting SNP TSC0014835.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABC53910;
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                                                                                                                                                                                                                                                                                                                                       EPIGENOMICS
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                                                                                                                                                                                                                                                                               Piepenbrock C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TGTTTAGGGAAGA 13
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Pred. No. 89;
                                                                                                                                      for diagnosis and cell otide polymorphisms and
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This invention describes novel oligonucleotide primers or peptide nucleic

Claim 1; SEQ ID NO 53927; 29pp + Sequence Listing;

German

Sequence 13 BP; 3 A; 6 C; 0 G; 4 T; 0 U; 0 Other;

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RESULT 149
ABPO7049/c
ID ABPO70
XX ABPO70
XX ABPO70
XX SNP; B
XW Peptid
XW Centra
XX SNP; B
XW WO2001
XX WO2001
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XX O1-APR
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                                                                                                                         This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99389, ABF00010-ABF99389, ABF000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 89;
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RESULT 150
ABF18399/c
ID ABF183
XX ABF183
XX ABF183
XX ABF183
XX PAFEB
XX Oligon
XX SNP; s
XW Peptid
XW Centra
XX Homo s
XX WO2001
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RESULT 151
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ID ABF888
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   ABF88832
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; SEQ ID NO 118396; 29pp + Sequence Listing; German.
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nilarity 84.6%;
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                                                                                                                                                                                                                                                                                                                                                                                central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9989, ABF00010-ABF99989, ABH00010-ABH00910-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; peptide nucleic acid; cytosine methylation; cardiovascular; primer; central nervous system; gastrointestinal; respiratory; immune; metal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                               Oligonucleotide
                                                                                                                           22-FEB-2002
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                                                                                                                                                                                  ABH64890 standard; DNA;
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central nervous
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                         Olek A,
                                                                              07-APR-2000; 2000DE-01019173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                  Sequence 13 BP; 4 A; 0 C; 7 G; 2 T; 0 U; 0 Other;
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  1725 TTGAGGGAACAGA 1737
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                                                                   20.4%;
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Pred. No. 89;
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RESULT 154
ABC53911/c
ID ABC539
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KW Peptic
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                                                                                                                                                                                                                                                                                                                                                                                                             respiratory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TSC0014835
                                                                                                                                                                                                                                                                                                                                                                                                             immune;
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The

Set of oligonucleotides, designed to detect singlemethylation status.

nnucleotides, useful for diagnosis and cell typing, detect single-nucleotide polymorphisms and cytosine

WPI; 2001-657177/75.

Olek A,

Piepenbrock C,

Berlin

ζ,

07-APR-2000; 2000DE-01019173 06-APR-2001; 2001WO-IB000713.

EPIGENOMICS

ĀG

Claim 1; SEQ ID NO 53928; 29pp + Sequence Listing; German

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RESULT 155
ABC83524
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       oligomers are also used for detecting cell type differentiation. ABC000 -ABC99989, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF3997073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                      acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
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                                                                                                                                                                                                                                                                                                            methylation status.
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 Conservative
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Pred. No. 89;
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Mismatches
                                                      0 U;
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RESULT 157
ABF11987
ID ABF119
XX
AC ABF119
XX
DT 21-FEB
XX

ABF11987

standard;

DNA;

13

ВP

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21-FEB-2002

(first entry)

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RESULT 156
ABF11986/c
ID ABF119
XX ABF119
XX ABF119
XX Oligon
XX SNP; B
KW Peptid
XW Centra
XX Homo B
XX H
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention describes novel oligonucleotide primers or peptide nucleic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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AGAGAGGAGAATT 13
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RESULT 158
ABH10409/c
ID ABH10409;
XX
AC ABH10409;
XX
Oligonucleotide SEQ ID NO 210
XX
SNP; single nucleotide golymc
KW septide nucleic acid; cytosir
KW peptide nucleic acid; cytosir
KW central nervous system; gastr
XX
PN W0200177384-A2.
XX
PF 06-APR-2001; 2001WO-IB000713.
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   06-APR-2001; 2001WO-IB000713.
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                                                                                       system; gastrointestinal;
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Pred. No. 89;
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                                                                                                                                  detecting SNP
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                                                                                        respiratory; immune; metabolic
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RESULT 159
ABH44535/c
ID ABH445
XX ABH445
XX ABH445
XX SNP; s
XW SNP; s
XW peptid
XW Centra
XX Homo s
XX Homo s
XX HOTO s
XX H
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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF0010-ABH99989, ABH0010-ABH99989 and ABI00010-ABH99989, represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
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                                                                                                                                                                                                                                                                                                          Set of oligonuclectides, useful for diagnosis and cell typing, designed to detect single-nuclectide polymorphisms and cytosine
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                                                                                                                                                                                                           This invention describes novel oligonucleotide primers or peptide nucleic
                                                                                                                                                                                                                                                 Claim 1; SEQ ID NO 250250; 29pp + Sequence Listing; German
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                                                                                                                                                                                                                                This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; car peptide nucleic acid; cytosine methylation; cardiovascular; pricentral nervous system; gastrointestinal; respiratory; immune;
                                                                                                                                                                                   Sequence 13
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                                                                                           Local Similarity
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Pred. No. 89;
                                                                                              Score 9.8; D
Pred. No. 89;
                                                                     Mismatches
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RESULT 163
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                Oligonucleotide SEQ ID NO 74892 for detecting SNP TSC0019229.
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                                                                                                      21-FEB-2002
                                                                                                                                     ABC74875
                                                                                                                                                                   ABC74875 standard; DNA; 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Olek A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-APR-2000; 2000DE-01019173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide SEQ ID NO 72972 for detecting SNP TSC0018828.
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                                                                                                                                                                                                                                                                                  1736 GACAGGAGAAATG 1748
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                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                    (first entry)
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RESULT 164
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Best Local
                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; peptide nucleic acid; cytosine methylation; cardiovascular; primer; central nervous system; gastrointestinal; respiratory; immune; metal
                                                                                                                                             Oligonucleotide SEQ
                       07-APR-2000; 2000DE-01019173.
                                                                                                                                                                                              ABC10735 standard; DNA; 13
                                                                                                                                                                                                                                                                                                            Sequence 13 BP; 6 A; 5 C; 0 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosin methylation status.
                                        06-APR-2001; 2001WO-IB000713.
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                                                                                                                                                                              ABC10735
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1; SEQ ID NO 74892; 29pp + Sequence Listing; German
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                                                                                                                                                                                                                                                         1718 ACTGATGTTGAGG 1730
                                                                                         sapiens.
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                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                           ID NO 10726 for detecting SNP TSC0002683.
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Pred. No. 89;
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(EPIG-) EPIGENOMICS

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RESULT 165
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XX ABC124
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XX SNP; BE
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KW Centra
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Best Local S
Matches 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                    Set of oligonucleotides, useful for designed to detect single-nucleotide methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                  Claim 1; SEQ ID NO 12433;
                                                                                                                                                                                                                                                                                          WPI; 2001-657177/75
                                                                                                                                                                                                                                                                                                                                            Olek A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oligonucleotide SEQ ID NO 12433 for detecting SNP TSC0002943.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-APR-2000; 2000DE-01019173
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                                                                                                                                                                                                                                                                                                                                                                                                       (EPIG-)
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                                                                                                                                                                                                                                                                                                                                            Piepenbrock C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATAAAGGAGAAAT 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                            Berlin
                                                                                                               29pp + Sequence Listing;
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Pred. No. 8
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Mismatches
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                                                                                                                                                                                                  polymorphisms
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The

Sequence 13

BP; 4 A; 4 C; 0 G; 5 T; 0 U;

0 Other; DB 1:

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RESULT 166
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XX ABF023
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XX Oligon
XX SNP; s
KW Peptid
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XX H
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Best Local (
                                                                       range of diseases including immune system, gastrointestinal, respirator central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC000-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosin methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 102352; 29pp + Sequence Listing; German.
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                                                      ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     system; gastrointestinal; respiratory; immune;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 entry)
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Pred. No. 89;
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cardiovascular; primer;
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RESULT 168
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                                                                                                                                                                  Query Match
Best Local Similarity
Matches 11; Conserv
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Matches 11; Conservative
                                                                                                                                                                                                                                                     This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                              ABF46517 standard; DNA; 13 BP
                                                                                                                                                                                                                             Sequence 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; SEQ ID NO 114028; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-657177/75
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (EPIG-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                      1744 AAATGCATCCATT 1756
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                                                                                                                                                                                                                             BP;
                                                                                                                                                                    Conservative
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                                                                                                                                                                                 20.4%;
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                                                                                                                                                                 Score 9.8; DB Pred. No. 89; O; Mismatches
                                                                                                                                                                    0;
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0; Mismatches
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ABF46517;

US6559125-B1

Human immunodeficiency virus

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ADD15389/c
ID ADD153
XX
AC ADD153
XX
DT 15-JAN
XX
Plasmi
DE Plasmi
XX
KW ss; po
KW ss; po
KW knocko
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Uniden
OS Uniden
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PN US6559
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                                                                                                                                                                                                                                                                                                                                                             Matches
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABI32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosin methylation status.
                                                           Synthetic
                                                                                                                                    Plasmid pHIV-LTR EcoRI/HindIII restriction fragment 2 with target DNA
                                                                                                                                                                  15-JAN-2004
                                                                                                                                                                                               ADD15389
                                                                                                                                                                                                                          ADD15389 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 13 BP; 3 A; 4 C; 0 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; SEQ ID NO 146514; 29pp + Sequence Listing; German
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-APR-2000; 2000DE-01019173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oligonucleotide SEQ ID NO 146514 for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-FEB-2002
                                                                                       knockout;
                                              Inidentified.
                                                                                          polyamide alkylator; conjugate; hairpin; regulator; gene therapy;
ckout; pHIV-LTR EcoRI/HindIII.
                                                                                                                                                                                                                                                                                                                              1721 GATGTTGAGGGAA 1733
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                                                                                                                                                                                                                                                                                                   GATGTTGAAAGAA 1
                                                                                        PHIV-LTR
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                                                                                                                                                                                                                          ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                diagnosis;
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ARESULT 170
ABT39656/c
ID ABT39658/c
AC ABT396
XX ABT396
XX T12-JUN
XX T12-JUN
XX CYtost
KW CYtost
KW Schizo
KW Schizo
KW Homo S
XX WO2003
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XX MOLE-
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention relates to a novel polyamide alkylator conjugate. Specifically, it refers to a hairpin polyamide comprising a pyrrole and/ or imidazole amino acid linked to a gamma aminobutyric acid, which in turn is linked to the alkylator that selectively alkylates only one strand of a double-stranded DNA molecule. The present invention describes a conjugate that can be used to target a predetermined DNA sequence and thereby inhibit DNA-protein interactions, and hence provides a novel regulator of gene expression. As such, in addition to competing with transcription factors, the conjugates can be used in gene therapy to target a gene's coding region for use as a knockout reagent. This oligonucleotide sequence is restriction fragement 2 derived from the plasmid pHIV-LTR ECORI/HindIII that contains the target DNA sequence of the oliverties.
   WPI; 2003-313353/30
                                                                                                                                                                                                                                                         schizophrenia;
human fukutin;
                                                                                                                                                                                                                                                                                         Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease;
                               Telerman A,
                                                                                                                              17-SEP-2002; 2002WO-IB004208.
                                                                                                                                                               27-MAR-2003.
                                                                                                                                                                                             WO2003025175-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                        ABT39656 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Polyimide-alkylator conjugate for therapeutic purposes, comprises alkylator linked to polyamide having pyrrole and/or imidazole amino acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-JAN-2001; 2001US-00772315.
                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                         Tumour suppression related human fukutin oligo
                                                                                                                                                                                                                                                                                                                                                                       12-JUN-2003 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                         ABT39656;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 13 BP; 2 A; 4 C; 4 G; 3 T; 0 U; 0 Other;
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                                                                (MOLE-) MOLECULAR ENGINES LAB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the polyamide alkylator conjugate of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1741 GAGAAATGCATCC 1753
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                                 Amson R,
                                                                                                2001FR-00011978.
                                                                                                                                                                                                                                                             protein chip; gene ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20.4%;
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                                 Tuijnder
                                                                                                                                                                                                                                                                                                                                                                                                                                        ВP
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Pred. No. 89;
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                                                                                                                                                                                                                                                                          therapy; tumour suppression;
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XFFFX8XSSSSSSSSSSSSSSSSSSSSSSSSSSS New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.

Disclosure; Page 652; 720pp; French.

The invention relates to a novel isolated 17 mer nucleic acid sequence, CC given in the specification, a sequence containing at least 15 consecutive CC nucleotides from the 17 mer sequence, a sequence with, after optimal CC alignment, at least 80 % identity to the 17 mer sequence, a sequence that CC of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, CC identifying, quantifying and/or amplifying a nucleic acid, e.g. as one CC component of a gene chip, in vitro as (anti) sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, collapped for containing the nucleic acids, cells containing the cyector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral CC diseases that are characterised by development of tumours or cell CC diseases that are characterised by development of tumours or cell CC diseases. The polypeptides can also be used to generate antibodies, and coth the polypeptide and antibodies are useful as components of these CC diseases. The polypeptides can also be used to generate antibodies, and coth the polypeptide and antibodies are useful as components of protein CC chips. The nucleic acid sequences of the invention can be used in gene CC related human fukutin oligonucleotide of the invention as tumour suppression

Sequence 17 BP; 8 A; 2 C; 6 G; 1 T; 0 U; 0 Other;

Query Match Best Local ( 10; Similarity Conservative 18.3%; <u>,</u> Score 8.8; DB 1; Pred. No. 1.3e+02 Mismatches DB 1; Length 17; Indels 0; Gaps

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Search completed: July 13, 2004, 11:03:43 Job time : 1 secs

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Maximum DB
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Maximum Match
Listing first
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Perfect score:
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Match
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Gapop 10.0 , Gapext 0.5
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US-09-252-292-27

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US-09-866-108A-8345

US-09-866-108A-8345

US-09-866-108A-8345

US-09-866-108A-8345

US-09-866-108A-8345

US-09-861-108A-8345

US-09-861-108A-8345

US-09-81-46-601

US-08-29-620A-78

US-09-091-46-601

US-08-158-071-8

US-08-284-746-17

US-08-488-551B-516

US-08-488-551B-516

US-08-488-551B-834

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Sequence 27, Appl Sequence 23, Appl Sequence 341, Ap Sequence 8341, Ap Sequence 8344, Ap Sequence 8343, Ap Sequence 8344, Ap Sequence 8345, Appl Sequence 5, Appl Sequence 5, Appl Sequence 78, Appl Sequence 78, Appl Sequence 78, Appl Sequence 8, Appli Sequence 8, Appli Sequence 10, Appl Sequence 11, Appl Sequence 515, Appl Sequence 516, Appl Sequence 516, Appl Sequence 517, Appl Sequence 518, Appl Sequence 516, Appl Sequence 517, Appl Sequence 518, Appl Sequence 517, Appl Sequence 518, Appl Sequence 517, Appl Sequence 518, Appl Sequence 517, Appl Sequence 518, Appl Sequence 518, Appl Sequence 519, Appl 
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Query Match 30.8%; Score 14.8; DB 1; Length 18; Best Local Similarity 88.9%; Pred. No. 2.9; Matches 16; Conservative 0; Mismatches 2; Indels 0; Gaps Qy 1714 GCTGACTGATGTTGAGGG 1731                    Db 18 GCTGACTGACGCTGAGGG 1	US-08-997-340-27/c  SEQUENCE 27, Application US/08897340  Patent No. 595306  Patent No. 595306  Patent No. 595306  Patent No. 595306  REPLICATION: Gimeno, Carlos J. and Errada, Patrick, R. TITLE OF INVENTION: Weight Control Pathway Genes and Uses TITLE OF INVENTION: Therefor NUMBER OF SEQUENCES: 36  CORRESPONDENCE ADDRESS: ADDRESSE: LAHIVE & COCKFIELD, LLP STREET: 28 State Street  CITY: Boston  SYNTE: Massachusetts  COUNTRY: USA  ZIP: 02109  COMPUTER READABLE FORM: REDUCKET STREET: USA  COMPUTER READABLE FORM: Compatible OPERATING SYSTEM: PC.DOS/MS-DOS SOOTWARE: PatentIn Release #1.0, Version #1.25  CURRENT APPLICATION NUMBER: US/08/MS-DOS SOOTWARE: PATENCATION: 435  PRIOR APPLICATION NUMBER: US/08/897,340  FILLING DATE: 17-SEP-1996  ATTORNAY/AGENT INFORMATION: RECESTRATION NUMBER: US/08/897,340  RECISTRATION NUMBER: US/08/897,340  RECISTRATION NUMBER: US/08/897,340  TELEPHONE: (617)227-7400  TELEPHONE: (617)227-7400  TELEPHONE: (617)227-7400  TELEPHONE: (617)227-7400  TELEPHONE: GENERACTERISTICS: LENGTH: 18 base pairs  LENGTH: 18 base pairs  MOLECULE TYPE: CDNA  US-08-897-340-27	34 9 18.8 11 1 US-08-192-942-8 Sequence 8, Appli 35. 9 18.8 11 1 US-08-646-695-15 Sequence 15, Appl 1 0S-08-646-695-15 Sequence 15, Appl 2 18.8 12 1 US-08-173-489C-256 Sequence 25.6, Appl 38 9 18.8 12 1 US-08-173-489C-256 Sequence 25.6, Appl 2 18.8 12 1 US-08-6507-032-14 Sequence 14, Appl 2 18.8 12 1 US-08-6244-087-12 Sequence 12, Appl 2 18.8 12 1 US-08-244-087-12 Sequence 12, Appl 3 18.4 15.4 15.1 US-09-866-108A-8341 Sequence 612, Appl 3 18.6 17 1 US-09-866-108A-8342 Sequence 8341, Appl 3 18.6 17 1 US-09-866-108A-8342 Sequence 8342, Appl 3 18.6 12.1 10 1 US-09-866-108A-8343 Sequence 8343, Appl 3 18.6 12.1 10 1 US-09-866-108A-8344 Sequence 8343, Appl 3 18.6 12.1 10 1 US-08-866-108A-8345 Sequence 8343, Appl 3 18.6 12.1 10 1 US-08-866-108A-8345 Sequence 8343, Appl 5 18.8 12.1 10 1 US-08-866-108A-8345 Sequence 8343, Appl 5 18.8 12.1 10 1 US-08-488-551B-833 Sequence 515, Appl 5 18.8 12.1 10 1 US-08-488-551B-833 Sequence 8334, Appl 5 18.8 12.1 10 1 US-08-488-551B-833 Sequence 8334, Appl 5 18.8 12.1 10 1 US-08-488-551B-833 Sequence 8334, Appl 5 18.8 12.1 10 1 US-08-488-551B-833 Sequence 8334, Appl 5 18.8 12.1 10 1 US-08-488-551B-833 Sequence 8334, Appl 5 18.8 12.1 10 1 US-08-488-551B-833 Sequence 8334, Appl 5 18.8 12.1 10 1 US-08-488-551B-833 Sequence 8334, Appl 5 18.8 12.1 10 1 US-08-488-551B-833 Sequence 8334, Appl 5 18.8 12.1 10 1 US-08-488-551B-833 Sequence 8334, Appl 5 18.8 12.1 10 1 US-08-488-551B-833 Sequence 8334, Appl 5 18.8 12.1 10 1 US-08-488-551B-833 Sequence 8334, Appl 5 18.8 12.1 10 1 US-08-488-551B-833
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RESULT 2 US-09-252-329-27/c

Sequence 27, Application US/09252329 Patent No. 6147192 GENERAL INFORMATION:

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US-09-252-329-27
                                                                                                                                                                                                                                                    Sequence 23, Application US/09679298A Patent No. 6566131
GENERAL INFORMATION:
                                                                           SEQ ID NO 23
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                    CURRENT APPLICATION NUMBER: US/09/679,298A CURRENT FILING DATE: 2001-03-05 NUMBER OF SEQ ID NOS: 47
                                                                                                                                                                                APPLICANT: Brett P. Monia
APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF SMAD6 EXPRESSION
FILE REFERENCE: RTS-0045
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Silveri, Jean M.
REGISTRATION NUMBER: 39,030
REFERENCE/DOCKET NUMBER: MNI-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
                    ORGANISM: Artificial Sequence FEATURE:
                                                             TYPE: DNA
OTHER INFORMATION: Antisense Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOPTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/252,329
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Gimeno, Carlos J. and Errada, Patrick, R. TITLE OF INVENTION: Weight Control Pathway Genes and Uses TITLE OF INVENTION: Therefor NUMBER OF SEQUENCES: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: Massachusetts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                        1714 GCTGACTGATGTTGAGGG 1731
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Similarity 88.9%;
16; Conservative
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Pred. No. 2.
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GENERAL INFORMATION:
APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharre;
APPLICANT: HANZEL, David
APPLICANT: RANK, David
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Best Local Similarity
Matches 9; Conserva
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                                                           CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR PPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
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NUMBER OF SEQ ID NOS: 14225
SOFTWARE: PatentIn version 3.0
SEQ ID NO 5840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 8341, Application US/09866108A Patent No. 6686188
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Becobedo, Jaime
ITITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Rel
ITITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,876-J (237/198)
FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT FILING DATE: 1990-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-10-26
                                                                                                                                                                                                                                                               APPLICANT: HANZEL, David K.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 08/584,040
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 16
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
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o. 6566127
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Pred. No. 3
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Pred. No. 4.7;
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TITLE OF INVENTION: TITLE OF INVENTION: TITLE OF INVENTION: PILE REFERENCE: AEOMICA-7

FILE REFERENCE: AEOMICA-7

CURRENT APPLICATION NUMBER: US 60/207,456

PRIOR APPLICATION NUMBER: US 60/207,456

PRIOR FILING DATE: 2000-05-26

PRIOR APPLICATION NUMBER: GB 24263.6

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR PILING DATE: 2000-09-27

PRIOR PILING DATE: 2000-09-27
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US-09-866-108A-8341
                                                                                                                                                                                                PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
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SEQ ID NO 8341
LENGTH: 17
TYPE: DNA
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PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR PPLICATION NUMBER: PCT/US01/00665
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
                                                                     Remaining Prior Application data removed - NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Aeomica Sequence Listing Engine Patent No. 6686188
SEQ ID NO 8342
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SOFTWARE: Acomica Sequence Listing Engine
ORGANISM: Homo sapiens
                                                  ENGTH:
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PENN, Sharron G.
David K.
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; ORGANISM: Homo sapiens
US-09-866-108A-8343
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Patent No. 6686188
SEQ ID NO 8343
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                                                                   Matches
                                                                                 Query Match
Best Local Similarity
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
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PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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                       1713 TGCTGACTGATGT 1725
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15 TGCTGACTGATGT 3
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13; Conserv
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CHEN, Wenshen
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                                                                                 27.1%;
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100.0%; Pred. No.
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                                                                   Mismatches
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US-09-866-108A-8344/c ; Sequence 8344, Application US/09866108A ; Patent No. 6686188

GENERAL INFORMATION:
APPLICANT: GU, Yizhong
APPLICANT: JI, Yongga

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FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 60/2036,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron
APPLICANT: HANZEL, David
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
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Best Local
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SEQ ID NO 8344
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PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
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PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
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CURRENT FILING DATE: 2001-05-25
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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PRIOR FILING DATE: 2001-01-30
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TYPE: DNA
ORGANISM: Homo sapiens
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DR APPLICATION NUMBER: PCT/US01/00667

DR FILING DATE: 2001-01-30

DR FILING DATE: 2001-01-30

DR FILING DATE: 2001-01-30

DR APPLICATION NUMBER: PCT/US01/00669

DR FILING DATE: 2001-01-30

DR FILING DATE: 2001-01-30
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Local Similarity 100.0%;
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HANZEL, David K.
RANK, David R.
CHEN, Wensheng
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HANZEL, David
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Pred. No. 5.8;
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; Sequence 67, Application US/08666341A
; Patent No. 6365345
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                                                        US-08-666-341A-67
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NUMBER OF SEO ID NOS: 15755
SOFTWARE: Acomica Sequence Listing Engine
Patent No. 6686188
SEO ID NO 8345
 Matches
             Query Match
Best Local
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR TILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 17
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disc
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                     MOLECULE TYPE:
ANTI-SENSE: Y
                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                       CITY: Washington STATE: D.C.
                                                                                                              STRANDEDNESS:
                                                                                                                         TYPE: nucleic acid
                                                                                                                                                                                                                             APPLICATION NUMBER: FILING DATE: 15-AUC
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                                                                                                  TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
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              Similarity
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                                                                                                unknown
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                                                                                   DNA (genomic)
                                                                                                               unknown
             26.7%;
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prevention and treatment
of c-erbB plays a role
106
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                                                                                                                                                                                                                                                                     Release #1.0, Version
                                                                                                                                                                                   EP 93120710.4
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             Score 12.8;
Pred. No. 6;
  Mismatches
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5.8;
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disorders in which expression
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1723 TGTTGAGGGAACAGAC 1738

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Matches
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TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,876-0 (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOPTWARE: PatentIn version 3.0
SEQ ID NO 5841
LENGTH: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 5, Application Patent No. 5756294 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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Patent No. 6500___
Patent INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 5841, Application US/09371772B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
ATTORNEY/AGENT INFORMATION:
NAME: Swecker, Robert S.
REGISTRATION NUMBER: 19,885
REFERENCE/DOCKET NUMBER: 0201
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-836-6620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: Floppy disk
COMPUTER: PA PC compatible
COMPUTER: PA C compatible
COMPUTER: PA C compatible
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: US/08/533,472
FILING DATE: 25-SEP-1995
CLASSIFICATION: 435
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Sadzewicz, Lisa K.
TITLE OF INVENTION: Suceptibility Mutation for Breast and
TITLE OF INVENTION: Ovarian Cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: (
                                                                                                                                                                                                                                                                                                                                                                                STREET: 699 Prince CITY: Alexandria STATE: VA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity 62.9 10s 10; Conservative
                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
ZIP: 22314-3187
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Escobedo, Jaime
                                                                                                                                                                                                                                                                                                                                                                                                                              699 Prince Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              White, Marga B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 BURNS, DOANE, SWECKER & MATHIS
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62.5%; Pred. No. 6;
ative 4; Mismatches
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   TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base maintained.
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APPLICANT: Susan
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SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                 SOFTMARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION DATA:
                                                                                                    NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 200
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                 APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY/ACENT INFORMATION:
                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA: PRIOR APPLICATION DATA: PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
FILING DATE: August
CLASSIFICATION: 43:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                  TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: U.S.A. ZIP: 90071-2066
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
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nucleic acid
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Dan T. Stinchcom
James McSwiggen
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ENTION: RIBOZYME TREATMENT OF
ENTION: DISEASES OR CONDITIONS
TENTION: RELATED TO LEVELS OF
ENTION: INTRACELLULAR ADHESION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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N: 435
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                                                                                                                                                                                                                                                                                                                                                           US/08/292,620A
                                                                                                                                                                                                                                                                         including application described below:
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                                                                                                                                       208/149
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US-09-071-845-78/c
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TOPOLOGY:
US-09-071-845-78
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                    Query Match
Best Local Similarity
Matches 13; Conserv
                                                                                                                            INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pair
TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 78,
                                                                                                                                                                                                                                               FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208/14
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: RIBOZYME TREATMENT OF
TITLE OF INVENTION: BLASTED OF LEVELS OF
TITLE OF INVENTION: RELATED TO LEVELS OF
TITLE OF INVENTION: INTRACELLULAR ADHESION
TITLE OF INVENTION: MOLECULE-1 (I-CAM-1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/0
                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                  TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/292,6
FILING DATE: August 17, 1994
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: U.S.A.
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 1724 GTTGAGGGAACAGAC 1738
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6132967
                                  Conservative
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James McSwiggen
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Dan T. Stind
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                                                                                                                   linear
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                                                24.6%;
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                                  0;
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Pred. No. 8
                                              Score 11.8;
Pred. No. 8.
                                                                                                                                                                                                                                                                                   208/149
                                Mismatches
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                                                                 DB 1; Length 15;
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                                  Indels
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                                  Gaps
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TITLE OF INVENTION: Gene Expression Profiles in No. 6333152mal and TITLE OF INVENTION: Cancer Cells
FILE REFERENCE: 01107,74664
CURRENT APPLICATION NUMBER: US/09/081,646
CURRENT FILING DATE: 1998-05-20
EARLIER APPLICATION NUMBER: 60/047,352
EARLIER FILING DATE: 1997-05-21
NUMBER OF SEQ ID NOS: 871
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 601
LENGTH: 15
TYPE: DNA
ORGANISM: Homo sapiens
US-09-081-646-601
COMPUTER: IBM PC COMPACTIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/368,071
FILING DATE: 03-JAN-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: CAMPBELL, CATHRYN
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-LJ 1275
TELECOMMUNICATION INFORMATION:
TELEPAX: 619-535-9901
TELEPAX: 619-535-9901
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 base pairs
TYDE: "ATTORNES PAIRS
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US-08-368-071-8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: MILLAN, JO
TITLE OF INVENTION: I
TITLE OF INVENTION: I
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 8, Application US/08368071 Patent No. 5707853
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 601, Application US/09081646 Patent No. 6333152
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Best Local Similarity
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APPLICANT: Vogelstein, Bert
APPLICANT: Zhang, Lin
APPLICANT: Zhou, Wei
                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: UNITED STATES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: SAN DIEGO
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4370 LA JOLLA VILLAGE DRIVE, SUITE 700
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Pred. No. 10
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US-08-458-181-8
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                                                                                                                        Sequence 8, Application PC/TUS9302172 GENERAL INFORMATION:
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Best Local 9
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GENERAL INFORMATION:
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Best Local Similarity 91.7%;
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: CAMEBELL, CATHRYN
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-I
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/458,181
FILING DATE: 30-DEC-1994
CLASSIFICATION: 435
CLASSIFICATION: 435
CORRESPONDENCE ADDRESS:
ADDRESSEE: La Jolla Cancer Research Foundation STREET: 10901 North Torry Pines Road
                                                      APPLICANT: La Jolla Cancer Research Foundation
TITLE OF INVENTION: RECOMBINANT CALF INTESTINAL ALKALINE
TITLE OF INVENTION: PHOSPHATASE
NUMBER OF SEQUENCES: 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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ADDRESSEE: CAMPBELL AND FLORES
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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TYPE: nucleic acid
STRANDEDNESS: single
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Pred. No. 1:
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Pred. No. 1;
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US-08-173-489C-244/c
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                    FILING LALLS.

CLASSIFICATION: 435

PRIOR APPLICATION DATA:

APPLICATION UNMBER: US 07/968,436

FILING DATE: 29 OCT 1992

ATTORNEY/AGENT INFORMATION:

NAME: Handelman, Joseph H.

REGISTRATION NUMBER: U9518-6

REFERENCE/DOCKET NUMBER: U9518-6

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION (212) 708-1880
                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: GENETIC SEQUENCE ASSAY USIN TITLE OF INVENTION: TRIPLE-STRAND FORMATION.

NUMBER OF SEQUENCES: 365
CORRESPONDENCE ADDRESS:
ADDRESSEE: PROFILE DIAGNOSTIC SCIENCES, INC., STREET: 510 EAST 73RD STREET,
CITY: NEW YORK
STATE: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 2 Patent No.
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Best Local Similarity
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                                                                                                                                                                                                                                          ZIP: 10021.

COMPUTER READABLE FORM:
MEDIDY TYPE: 3.5 inch, 1.44Mb storage
COMPUTER: IBM PC/XT/AT
OPERATING SYSTEM: MS-DOS version 6.2
SOFTWARE: Wordperfect Version 5.1
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM CC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version 1.25
CURRENT APPLICATION DATA;
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LENGTH: 12 base pairs
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APPLICATION NUMBER: US/07/849,219
FILING DATE: 10-MAR-1992
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                                                                                                                                                                                                       APPLICATION NUMBER: US/0 FILING DATE: 22 DEC 1993
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STATE: California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1733 ACAGACAGGAGA 1744
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o. 5861244
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(attorney) (212) 246-8959
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENETIC SEQUENCE ASSAY USING DNA TRIPLE-STRAND FORMATION.
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Pred. No. 12;
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INFORMATION FOR SEQ ID NO:

ACTGGAGAAAGGC 13

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US-08-284-746-10
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                                                                                                         Matches
                                                                                                                                                                  Query Match
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GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/883,
FILING DATE: May 14, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
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ANTI-SENSE: no
PUBLICATION INFORMATION:
RELEVANT RESIDUES IN SEQ ID NO: 244 :FROM 1 TO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: James A. McSwiggen
TITLE OF INVENTION: ASSAY FOR RIBOZYME TARGET SITE
NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: other nucleic acid
DESCRIPTION: third strand derived from L.
DESCRIPTION: interrogans 23s region in Seq ID No. 5861244243
                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (213) 489-1600
                                                                                                                                         Local
                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 197/070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM compatible
OPERATING SYSTEM: IBM P.C. DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
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STRANDEDNESS: single stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
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1737 ACAGGAGAAATGC 1749
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                                                                                                                                         00.4%;
Similarity 84.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (213) 955-0440
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                                                                                                                                         Score 9.8;
Pred. No. 1
                                                                                                         Mismatches
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                                           Sequence 3, Application US/09772315

Patent No. 6559125

Patent No. 6559125

Patent No. 6559125

PAPPLICANT: DERVAN, Peter
APPLICANT: WUNTZ, Nicholas
APPLICANT: CHANG, Aileen
TITLE OF INVENTION: POLYAMIDE-ALKYLATOR CONJUGATES & RELATED PRODUCTS & METHODS
PILE REFERENCE: GENESOFT09/772315

CURRENT APPLICATION NUMBER: US/09/772,315

CURRENT PILING DATE: 2001-01-26

NUMBER OF SEQ ID NOS: 25

SOFTWARE: Patentin version 3.0

SEQ ID NO 3
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO:
               TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: May 14, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 197/070
                                  ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 435
PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (213) 955-0440
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OPERATING SYSTEM: IBM P.C. DOS (Version 5.0)
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611 West Sixth
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ASSAY FOR RIBOZYME TARGET SITE
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Pred. No. 17
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US-08-025-038-11/c
                                                                                          US-08-545-785-2/
                                                                                                              RESULT 24
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                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (414)289-3791
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 base pairs
TYPE: NUCLEIC ACID
                                  Sequence 2, Application US/08545785
Patent No. 5770713
GENERAL INFORMATION:
                                                                                                                                                                                                                                  Query Match 19.6%;
Best Local Similarity 90.9%;
Matches 10; Conservative
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Best Local (
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FILING DATE: 19930301
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/544,218
FILING DATE: 27-JUN-1990
ATTORNEY/AGENT INFORMATION:
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APPLICANT: BAXTER
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OTHER INFORMATION: Description of Artificial Sequence: pHIV-LTR EcoRI/HindIII
OTHER INFORMATION: restriction fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Meyers, Philip G.
REGISTRATION NUMBER: 30,478
REFERENCE/DOCKET NUMBER: 20
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy
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APPLICANT: Imbach and Rayner
TITLE OF INVENTION: Phosphorothioate Triester Oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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Local Similarity 84.6%;
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ZIP: 53202-5367
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CITY: Milwaukee
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Pred. No. 17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 515, Application US/08388353 Patent No. 6010895
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Best Local Similarity
Matches 10; Conserv
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CURRENT APPLICATION NUMBER: US/08/545,785
APPLICATION NUMBER: US/08/545,785
FILING DATE: 17-JAN 1996
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: JOSEPH LUCCI
REGISTRATION NUMBER: 33,307
REFERENCE/DOCKET NUMBER: ISIS-2114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
APPLICATION NUMBER: US/08/
FILING DATE: 14-FEB-1995
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: DIGIGILO, Frank S.
REGISTRATION NUMBER: 31,34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: And Method Of Preparation
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSE: Woodcock Washburn Kurtz Mackiewicz & No. 5770713ris LLP
STREET: One Liberty Place - 46th Floor
                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Scully, Scott, Murphy & Presser
STREET: 400 Garden City Plaza
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MEDIUM TYPE: 3.5 inch disk, 1.44 Mb
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 6.1
                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Cooper, David TITLE OF INVENTION: NON-
                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                       CITY: Garden City
STATE: New York
COUNTRY: United States
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                                                                                                                                                                                                                       ZIP: 11530
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STRANDEDNESS: single
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Jennifer
                                                                                           US/08/388,353
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Pred. No. 1
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                                                                                                                                  Version #1.25
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                Query Match
Best Local Similarity
Watches 9; Conserve
                                                                                          ; STRANDEDNESS: single TOPOLOGY: linear MOLECULE TYPE: DNA (genomic) US-08-388-353-516
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                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: DiGIGIO, Frank S.
REGISTRATION NUMBER: 31,346
REFERENCE/DOCKET NUMBER: 9606
TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 742-4343
TELEPHAX: (516) 742-4366
TELEPHAX: (230 901 SANS UR
INFORMATION FOR SEQ ID NO: 516:
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TELEFAX: (516) 742-4366
TELEX: 230 901 SANS UR
INFORMATION FOR SEQ ID NO: 519
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/388,353
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STATE: New York
STATE: "Thited St
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TITLE OF INVENTION: NON-PATHOGENIC STRAINS OF HIV-1
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                                                                                                                                                     TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                             FILING DATE: 1:
CLASSIFICATION:
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                                                                                                                                                                     LENGTH:
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1749 CATCCATTC 1757
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                             18.8%; Score 9; DB inlarity 100.0%; Pred. No. 18. Conservative 0; Mismatches
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Crowe, Suzanne
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100.0%; Pred. No
tive 0; Mismat
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Mismatches
                                             DB 1;
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                                                                                    Sequence 516, Application US/08488551B Patent No. 6015661
                   GENERAL INFORMATION:
APPLICANT: Nichola
APPLICANT: Dale A.
APPLICANT: David C
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APPLICANT: Nicholas J. Deacon
APPLICANT: Dale A. McPhee
                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (516) 742-4366 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC COMPUTED: COMPUTED: COMPUTED: COMPUTED: PATENTIAL SYSTEM: PC DOS/MS-DOS
SOPTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: 1
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 10 base pairs
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NAME: FRANK S. DIGIGLIO
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION: TELEPHONE: (516) 742-4343
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 23-DEC-1994
APPLICATION UNBER: US 08/
FILING DATE: 14-FEB-1995
APPLICATION NUMBER: PN3021
FILING DATE: 17-MAY-1995
FILING DATE: 17-MAY-1995
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NUMBER OF SEQUENCES:
       TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
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o. 6015661
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Dale A. McPhee
David Cooper
VENTION: NON-PATHOGENIC STRAINS OF HIV-1
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                                                    Nicholas J. Deacon
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100.0%; Fi
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100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PN0284 (AU)
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                                                                                                                                                                                                                                                                                            Length 10;
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                  APPLICANT:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: SCULLY, SCOTT, MURPHY & PRESSER
STREET: 400 GARDEN CITY PLAZA
CITY: GARDEN CITY
STATE: NEW YORK
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 742-4343
TELEFAX: (516) 742-4366
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 14-FEB-1995
APPLICATION NUMBER: PN3021/95
FILING DATE: 17-MAY-1995
ATTORNEY/AGENT INFORMATION:
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FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA: PM3864 (AU)
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CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                             TITLE OF INVENTION: NON-PATHOGENIC STRAINS OF HIV-1 NUMBER OF SEQUENCES: 841
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FILING DATE: 14-FEB
APPLICATION NUMBER:
FILING DATE: 21-FEB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: PN0284 (AU) FILING DATE: 23-DEC-1994 APPLICATION NUMBER: US 08/388,3
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                                                                                                    COUNTRY: U.S.A. ZIP: 11530-0299
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9; Conservative 0;
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                                                                                                                                                                                                                                                                                                         Nicholas J. Deacon
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                                                                                                                                                                                                                                                                  Dale A. McPhee
David Cooper
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; Pred. No. 18
0; Mismatches
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Best Local Similarity
Watches 9; Conserva
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Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
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TELEPHONE: (516) 742-4343
TELEFAX: (516) 742-4366
INFORMATION FOR SEQ ID NO: 833:
                                                                                                                                                                                      ZIP: 11530-0299

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Dale A. McPhee
APPLICANT: David Cooper
TITLE OF INVENTION: NON-PATHOGENIC STRAINS OF HIV-1
NUMBER OF SEQUENCES: 841
CORRESPONDENCE ADDRESS:
                                                                                                                                APPLICATION NUMBER: US/01
FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 17-MAY-1995
ATTORNEY/AGENT INFORMATION:
NAME: FRANK S. DIGIGLIO
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Nicholas J. Deacon
APPLICANT: Dale A. McPhee
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                                                                                                                                                                                                                                                                                                                                                STREET: 400 GARDE
CITY: GARDEN CITY
                                                                APPLICATION NUMBER: PM38
FILING DATE: 14-FEB-1994
APPLICATION NUMBER: PM401
FILING DATE: 21-FEB-1994
                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 10 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 14-FEB-1995
APPLICATION NUMBER: PN3021/95
FILING DATE: 17-MAY-1995
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                APPLICATION NUMBER:
                                                 APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
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5. 6015661
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       MBER: US 08/388,353
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14-FEB-1995
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100.0%; Pred. No.
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                                                                                                                  PM3864 (AU)
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                                                 PN0284 (AU)
                                                                                  PM4002 (AU)
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TELEPHONE: (516) 742-4343
TELEPAX: (516) 742-4366
INFORMATION FOR SEQ ID NO: 834:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 base pairs
TYPE: nucleic acid
                                                                                                                     TELEFAX: 650-903-3501
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17, Application US/08618834C
Patent No. 6361937
GENERAL INFORMATION:
APPLICANT: Stryer, Lubert
TITLE OF INVENTION: Computer-Aided
TITLE OF INVENTION: Sequencing
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Best Local Similarity 100.0%;
Matches 9; Conservative C
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STRIE: LOS ALTOS
STATE: CA
STATE: CA
                                                                                                                                                                                                                                                         FILING DATE:
ATTORNBY/AGENT INFORMATION:
NAME: Ritter, Michael J.
REGISTRATION NUMBER: 36,653
REFERENCE/DOCKET NUMBER: AFI
                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 435
PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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NAME: FRANK S. DIGIGLIO
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0
                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/618,834C FILING DATE: 19-MAR-1996
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1720 TGATGTTGA 1728
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4906 El Camino Real, Suite 205
                                 Conservative
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650-903-3501
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                            18.8%; Score 9; DB 1
100.0%; Pred. No. 18;
tive 0; Mismatches
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                                              DB 1;
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Watches 9; Conserve
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; Sequence 21, Application US/09508753B
; Patent No. 6544736
; GENERAL INFORMATION:
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SEQUENCE CHARACTERISTICS:
LENGTH: 10 base Tropy
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APPLICANT: HITCHOF FUNAKI
APPLICANT: Eiji OHARA
APPLICANT: Masanori WATAHIKI
TITLE OF INVENTION: Method for Synthesizing cDNA from mRNA sample
FILE REFERENCE: 00162/HG
CURRENT APPLICATION NUMBER: US/09/508,753B
CURRENT FILING DATE: 2000-06-16
PRIOR APPLICATION NUMBER: JP 9/270324
PRIOR FILING DATE: 1997-09-18
                                                                                                                            APPLICANT: Akira SHIMAMOTO
APPLICANT: Yasuhiro FURUICHI
APPLICANT: Yuko SHIBATA
APPLICANT: Hiroko FUNAKI
APPLICANT: Eiji OHARA
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APPLICANT: Stryer
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ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
NAME: Ritter, Michael J.
REGISTRATION NUMBER: 36,653
REFERENCE/DOCKET NUMBER: AFFY
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-903-3500
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FILING DATE: 19-MAR-11
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Stryer, Lubert
TITLE OF INVENTION: Comput
TITLE OF INVENTION: Seque
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CITY: Los Altos
STATE: CA
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TOPOLOGY: 1i1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
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100.0%; Pred. No. 18;
tive 0; Mismatches
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Sequencing
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                                                       Matches
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SEQ ID NO 21
                                                                                   Query Match
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APPLICANT: JAMES D.
                                                                                                                                                                                                    TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 squence 8, Application US/08192942
stent No. 5989906
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ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Description of Artificial Sequence: Primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: IBM COMPATIBLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ENGTH: 10
                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 197/173
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 1489-1600
                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: IBM P.C. DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               JUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                       Local Similarity
tes 8; Conserv
                                                                                                                                                                                                                                   TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET:
                                                                                                                                                                           LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                         FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE:
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                           1740 GGAGAAATG 1748
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 GGAGAAAUG 9
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611 West Sixth Street
                                                                                                                                                                                                                                      (213) 955-0440
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                                                         Conservative
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                                                                     18.8%;
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; Pred. No. 18;
0; Mismatches
                                                                    Score 9;
Pred. No.
                                                                     DB 1;
20;
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RESULT 35

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RESULT 36
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC COMPATIBLE
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NATA:
APPLICATION NUMBER: US/08/646,695
FILING DATE: On Even Date Herewith
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 6523-008
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: 66141 PENNIE INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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                        ZIP: 10036-2711

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: ROSE, John K. TITLE OF INVENTION: RECORTITLE OF INVENTION: USES
                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: unknown
                                                                                                                                                                            CITY: New York
STATE: New York
                                                                                                                                                                                                                   STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-9741/8864
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STATE: New York
                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                      ADDRESSEE:
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             PCT/US96/06053
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RESULT 37
US-08-173-489C-256/c
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TELEX: 66111 PENNIE
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 11 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 100.0%; Pred. No. 20; Matches 9; Conservative 0; Mismatches
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                                                                                                                                                                                                                                       FILING DATE: 22 DEC 1993
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 07/968,436
FILING DATE: 29 OCT 1992
ATTORNEY,AGENT INFORMATION:
NAME: Handelman, Joseph H.
REGISTRATION UNMBER: 26,179
REFERENCE/DOCKET NUMBER: U9518-6
TELECOMMUNICATION INFORMATION:
                                                                                                                                                            TELEFAX: (attorney) (2
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: WANG, C.-G.
APPLICANT: HEPBURN, A. G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Wordperfect Version CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/173.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44Mb storage
COMPUTER: IBM PC/XT/AT
OPERATING SYSTEM: MS-DOS version 6.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: PROFILE DIAGNOSTIC SCIENCES, INC.,
STREET: 510 EAST 73RD STREET,
CITY: NEW YORK
STATE: NEW YORK
               MOLECULE TYPE: other nucleic acid
DESCRIPTION: third strand derived from M. luteus
DESCRIPTION: 23s region in Seq ID No. 5861244255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: GENETIC SEQUENCE ASSAY USING DNATITLE OF INVENTION: TRIPLE-STRAND FORMATION.

NUMBER OF SEQUENCES: 365
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                        TYPE: nucleic acid
STRANDEDNESS: single stranded
TOPOLOGY: linear
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                                                                                                                                                                                                                        TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 6523-009-228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Misrock, S. Leslie REGISTRATION NUMBER: 18,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: unknown
                                                                                                                                         ENGTH:
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                                                                                                                                           12 bases
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                                                                                                                                                                                 (attorney) (212) 708-1880 attorney) (212) 246-8959 SEQ ID NO: 256:
yes
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22 DEC 1993
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                                                Query Match
Best Local Similarity
Matches 9; Conserve
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Best Local Similarity
Matches 9; Conserve
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; PUBLICATION INFORMATION:
; RELEVANT RESIDUES IN S
US-08-173-489C-256
                                                                                                                                    ; MOLECULE TYPE: DNA (genomic) US-08-507-032-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-507-032-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                     FILING DATE:
APPLICATION NUMBER: US 07/749
AFILICATION NUMBER: US 07/749
FILING DATE: 22-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 5490
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 12 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                         TELEFAX: 415-326-2422
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/507,032
FILING DATE:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
ADDRESSEE: William M. Smith
STREET: One Market Plaza, Steuart Tower, Suite 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Flanagan, William A. APPLICANT: Crabtree, Gerald R. TITLE OF INVENTION: Screening M TITLE OF INVENTION: Agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: One Market E
                                                                                                                                                                                                                                                                                           TELEPHONE:
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                                1738 CAGGAGAAA 1746
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                                                                18.8%; Score 9; DB illarity 100.0%; Pred. No. 21 Conservative 0; Mismatches
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415-326-2422
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22-AUG-1991
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100.0%; Pred. No. 21
tive 0; Mismatches
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                                                                    0.
                                                                                                  Length 12;
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                                                                    Indels
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RESULT 39 US-08-862-431-22 ; Sequence 22, Application US/08862431

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US-08-244-087-12/c
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/862
FILING DATE: 23-MAY-1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Kim, Judith U.
REGISTRATION NUMBER: 40,679
REFERENCE/DOCKET NUMBER: 1669
TELEPHONE: (202) 371-2600
TELEPHONE: (202) 371-2600
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 12, Application US/08244087 Patent No. 6610294
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                        APPLICANT: Chess, Leonard
APPLICANT: Yellin, Michael J.
TITLE OF INVENTION: MURINE MONOCLONAL ANTIBODY (5c8)
TITLE OF INVENTION: RECOGNIZES A HUMAN GLYCOPROTEIN ON THE SURFACE OF
TITLE OF INVENTION: T-LYMPHOCYTES, COMPOSITIONS CONTAINING SAME AND METHODS
TITLE OF INVENTION: OF USE
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/244,087
                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Cooper & Dunham
STREET: 30 Rockefeller Plaza
CITY: New York
STRIE: New York
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TITLE OF INVENTION: ANTIOXIDANT RESPONSIVE ELEMENT
NUMBER OF SEQUENCES: 51
CORRESPONDENCE ADDRESS:
ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX P.L.L.C.
STREET: 1100 NEW YORK AVENUE, SUITE 600
CITY: WASHINGTON
                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid STRANDEDNESS: sing TOPOLOGY: linear
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                                                                                                                                                                                  COUNTRY: United States of America
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ZIP: 20005-3934
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COMPUTER: IBM POF Compatible
OPERATING SYSTEM: PC-DDS/MS-DOS
SOFTWAREN PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/09955
FILING DATE: 19921116
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 0575/39757-A-PCT
TELECOMOUNICATION.
TELEPHONE: (212) 977-9550
TELEX: 42253 COOP UI
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 base pairs
TYPE: NUCLEIC ACID
STRANDEDNESS: Single
TOOPLOGY: lines*
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PCT-US92-09955-12/c
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 977-9550
TELEPAX: (212) 664-0525
TELEX: 422523 COOP UI
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
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TITLE OF INVENTION: T-L'
TITLE OF INVENTION: USE
TITLE OF INVENTION: USE
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Chess, Leonard
APPLICANT: Yellin, Michael J.
APPLICANT: Yellin, Michael J.
TITLE OF INVENTION: MURINE MONOCLONAL ANTIBODY (5c8)
TITLE OF INVENTION: RECOGNIZES A HUMAN GLYCOPROTEIN ON THE SURFACE OF
TITLE OF INVENTION: T-LYMPHOCYTES, COMPOSITIONS CONTAINING SAME AND METHODS OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: CDNA HYPOTHETICAL: NO
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TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: United States of America ZIP: 10112
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Floppy disk
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Query Match
Best Local Similarity
Watches 8; Conserve
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US-09-081-646-601
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US-09-081-646-601
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PCT-US92-09955-12
         PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: UZ 60/236,359
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00664
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Kinzler, Kenneth
APPLICANT: Vogelstein, Bert
APPLICANT: Vogelstein, Bert
APPLICANT: Zhang, Lin
APPLICANT: Zhou, Wei
TITLE OF INVENTION: Gene Expression Profiles in No. 6333152mal and
TITLE OF INVENTION: Cancer Cells
FILE REFERENCE: 01107.74664
CURRENT FILING DATE: 1998-05-20
EARLIER APPLICATION NUMBER: 60/047,352
EARLIER APPLICATION NUMBER: 60/047,352
EARLIER FILING DATE: 1997-05-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 871
SOFTWARE: FRATSEQ for Windows Version 3.0
SEQ ID NO 601
LENGTH: 15
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Patent No. 6333152
                                                                                                                                                                          APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
                                                                                                                                                                                                                                                                                                                                           APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharror
APPLICANT: HANZEL, David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
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2001-01-30
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88.9%;
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Pred. No. 4
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o. 21;
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; ORGANISM: Homo sapiens
US-09-866-108A-8342
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NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Acomica Sequence Listing Engine
Patent No. 6886188
SEQ ID NO 8341
LENGTH: 17
                                                                           Patent No. 668
SEQ ID NO 8342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 8342, Application US/09866108A Patent No. 6686188
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Best Local
                                                                                                                                    Remaining Prior Application data removed - NUMBER OF SEQ ID NOS: 15755
                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: GU, Yizhong
APPLICANT: JI, Yongga
APPLICANT: PENN, Shar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: PCT/USO1/00669
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00665
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00668
                                                                                                                                                                             PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
                                                                                                                   SOFTWARE: Aeomica
                                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
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                                                      LENGTH:
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APPLICATION NUMBER: PCT/US01/00665
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APPLICATION NUMBER: PCT/US01/00663
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Pred. No.
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46
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Query Match

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Length 17;

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RESULT 45
US-09-866-108A-8343
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                                                                                                                         US-09-866-108A-8344
                                                                                                                                           RESULT 46
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
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Sequence 8344, Application US/09866108A
Patent No. 6686188
GENERAL INFORMATION:
APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PANN, Sharzon G.
APPLICANT: HANZEL, David K.
                                                                                                                                                                                                                                                                                   Query Match
Best Local :
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APPLICANT: JI, Yonggan
APPLICANT: PENN, Sharr
APPLICANT: HANZEL, Day
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                                                                                                                                                                                                                                                                                                                                    ORGANISM: Homo sapiens
-09-866-108A-8343
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: AEOMICA-7
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APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00668
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00663
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HANZEL, David
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Pred. No. 46;
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Pred. No.
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               PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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Patent NO. 6686188
GENERAL INFORMATION:
APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
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Matches
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APPLICANT: SHANNON, MARK
APPLICANT: SHANNON, MASK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
FILE REFERENCE: ABOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
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PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
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CURRENT FILING DATE: 2001-05-25
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
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8; Conser
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HANZEL, David
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PCT/US01/00664
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US-08-388-353-515
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US-09-866-108A-8345
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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Best Local Similarity
Thes 8; Conserva
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NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Acomica Sequence Listing Engine
Patent No. 6686188
SEQ ID NO 8345
LENGTH: 17
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GENERAL INFORMATION:
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                                                                                                                    TELEFAX: (516) 742-4360
TELEX: 230 901 SANS UR
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                  FILING DATE: 14-FEB-1995
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: DIGIGILO, Frank S.
REGISTRATION NUMBER: 31,34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 96
TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 742-4343
                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/388,353
                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                   SEQUENCE CHARACTERISTICS:
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                                                             TYPE: nucleic acid
                                                                                                                                      TELEPHONE: (516) 742-4366
                              STRANDEDNESS: si
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STREET: 40
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: New York
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McPhee, Dale A.
Crowe, Suzanne
                                                                                                                                                                                                                                                                                                                                                                                                                                                     United States
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                DNA (genomic)
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; TOPOLOGY: 1:
; MOLECULE TYPE:
US-08-488-551B-515
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Best Local Similarity
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                                                                                                                                                              Query Match
Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 14-FEB-1995
APPLICATION NUMBER: PN3021/95
FILING DATE: 17-MAY-1995
ATTORNEY/AGENT INFORMATION:
NAME: FRANK S. DIGIGLIO
REFERENCE/DOCKET NUMBER: 9606Z
TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 742-4343
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Nicholas J. Deacon
APPLICANT: Dale A. McPhee
                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (516) 742-43-6
TELEFAX: (516) 742-43-6
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FO-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 07-JUN-
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: SCULLY, S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: David C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: PM400 PILLING DATE: 21-PPD APPLICATION NUMBER: PM400 PM400
                                                                                                                        Local Similarity nes 7; Conserv
                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: GARDEN CITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 23-DEC-1994
APPLICATION NUMBER: US 08/388,353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                             ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1743 GAAATGCAT 1751
                                                             1743 GAAATGCAT 1751
1 GGAATGGAT 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                10 base pairs
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                                                                                                                               Conservative
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                                                                                                                                                          12.1%;
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                                                                                                                               Score 5.8; DB Pred. No. 56; 0; Mismatches
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Pred. No. 56;
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US-08-488-551B-833

Sequence 833, Application US/08488551B Patent No. 6015661

RESULT 50

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Page 19
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GENERAL IMPORMATION:
APPLICANT: Nai-Alas J. Deacon
APPLICANT: Dala A. McPhae
APPLICANT: Dala A. McPhae
APPLICANT: Dala Cooper
ITILE OF INVENTION: NON-PATHOGRAIC STRAINS OF HIV-1
NUMBER OF SEQUENCES: 41
INVERSE OF SEQUENCES: 42
INVERSE OF SEQUENCES: 43
INVERSE OF SEQUENCES: 44
INVERSE OF SEQUENCE
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Result
No.
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                                                                                               RESULT 1
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Maximum Match 100%
Listing first 20 summaries
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Maximum DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Perfect score:
Sequence 53, Application US/10000213
Publication No. US20030125271A1
GENERAL INFORMATION:
APPLICANT: Brenda F. Baker
APPLICANT: Mark P. Roach
APPLICANT: Kenneth Dobie
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score grea
and is der
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Match
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43.488 Million cell upda
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                 GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          using sw
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US-10-000-213-54

US-10-000-213-56

US-10-138-674-7673

US-10-287-949A-7673

US-10-327-805-23

US-10-327-805-23

US-10-327-805-23

US-09-740-332-3026

US-09-817-879-3026

US-10-138-674-7674

US-10-287-949A-7674

US-10-287-949A-5840

US-10-287-949A-8431

US-09-866-108-8343

US-10-138-674-5841
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Sequence 54, Appl
Sequence 56, Appl
Sequence 7673, Ap
Sequence 7673, Ap
Sequence 7673, Ap
Sequence 3026, Appl
Sequence 3026, Ap
Sequence 3026, Ap
Sequence 7674, Ap
Sequence 7674, Ap
Sequence 5840, Ap
Sequence 5840, Ap
Sequence 8341, Ap
Sequence 8342, Ap
Sequence 8342, Ap
Sequence 8343, Ap
Sequence 8343, Ap
Sequence 8344, Ap
Sequence 8341, Ap
Sequence 8341, Ap
Sequence 8341, Ap
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Sequence 55, Application US/10000213

Publication No. US20030125271A1

GENERAL INFORMATION:

APPLICANT: Brenda F. Baker

APPLICANT: Mark P. Roach

APPLICANT: Kenneth Dobie

TITLE OF INVENTION: ANTISENSE MODULATION OF VIT

FILE REFERENCE: RTS-0327

CURRENT APPLICATION NUMBER: US/10/000,213

CURRENT FILING DATE: 2001-11-14

NUMBER OF SEQ ID NOS: 94

SEQ ID NO 55

LENGTH: 20
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; ORGANISM: Artificial Sequence
; PEATURE:
; OTHER IMFORMATION: Antisense Oligonucleotide
US-10-000-213-53
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; OTHER INFORMATION: Antisense Oligonucleotide
US-10-000-213-54
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SEQ ID NO 54
LENGTH: 20
TYPE: DNA
ODGSSTORM
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Publication No. US20030125271A1

GENERAL INFORMATION:
APPLICANT: Brenda F. Baker
APPLICANT: Mark P. Roach
APPLICANT: Kenneth Dobie

TITLE OF INVENTION: ANTISENSE MODULATION OF VITAMIN D NUCLEAR RECEPTOR EXPRESSION
FILE REFERENCE: RTS-0327
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Best Local S
Matches 20
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CURRENT APPLICATION NUMBER: US/10/000,213
CURRENT FILING DATE: 2001-11-14
NUMBER OF SEQ ID NOS: 94
SEQ ID NO 53
LENGTH: 20
TYPE ON
                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/000,213
CURRENT FILING DATE: 2001-11-14
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Antisense Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence
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les 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                       1722 ATGTTGAGGGAACAGACAGG 1741
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100.0%; Pr/
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0; Mismatches
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3.2;
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3.2;
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RESULT 6
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                                                                                                                                                                                                    ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-138-674-7673
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  밁
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US-10-000-213-56
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                                                                                                                                                                                                                                                          APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00-876-N (400/049)
CURRENT APPLICATION NUMBER: US/10/138,674
CURRENT FILING DATE: 2002-05-03
NUMBER OF SEQ ID NOS: 20822
SOFTWARE: PatentIn version 3.0
SEQ ID NO 7673
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 7673, Application US/10138674 Publication No. US20040077565A1 GENERAL INFORMATION:
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SEQ ID NO 56
LENGTH: 20
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                                                                                                                                     Matches
                                                                                                                                                  Query Match
Best Local Similarity
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APPLICANT: Mark P. Roach
APPLICANT: Kenneth Dobie
TITLE OF INVENTION: ANTISENSE MODULATION OF VITAMIN D NUCLEAR RECEPTOR EXPRESSION
FILE REFERENCE: RTS-0327
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CURRENT FILING DATE: 2001-11-14
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ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                      LENGTH: 17
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                                                                                                   1715 CTGACTGATGTTGAGG 1730
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100.0%; Pr
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Pred. No. 9.1;
5; Mismatches
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; Pred. No. 3.2;
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Sequence 3026, Application US/09740332
Publication No. US20030125270A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
APPLICANT: RIDOZYme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Related TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: RPI 400/003
CURRENT APPLICATION NUMBER: US/09/740,332
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9704
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APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Pavco, Pam

APPLICANT: McSwiggen, Jim

APPLICANT: Stinchcomb, Dan

APPLICANT: Stinchcomb, Dan

APPLICANT: Stinchcomb, Dan

APPLICANT: Stinchcomb, Dan

APPLICANT: Besobedo, Jaime

TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions

TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor

FILE REFERENCE: MBHB00-876-N (400/049)

CURRENT APPLICATION NUMBER: US/10/287,949A

CURRENT FILING DATE: 2003-04-11

NUMBER OF SEQ ID NOS: 20822

SOFTWARE: Patentin version 3.0

SEQ ID NO 7673
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Publication No. US20030144241A1
GENERAL INFORMATION:
APPLICANT: Brett p. Monia
APPLICANT: Lex M. Cowsert
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SEQ ID NO 23
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                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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TITLE OF INVENTION: ANTISENSE MODULATION OF FILE REFERENCE: RTS-0045
CURRENT APPLICATION NUMBER: US/10/327,805
CURRENT FILING DATE: 2002-12-20
PRIOR APPLICATION NUMBER: US/09/679,298
PRIOR FILING DATE: 2001-03-05
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TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
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Pred. No. 9.
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                                            APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00-876-N (400/049)
CURRENT FILING DATE: 2002-05-03
NUMBER OF SEQ ID NOS: 20822
SOFTWARE: PatentIn version 3.0
SEQ ID NO 7674
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Ribozyme Pharmaceüticals inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: MBHB00-801-F
CURRENT APPLICATION NUMBER: US/09/817,879
CURRENT FILING DATE: 2001-03-26
NUMBER OF SEQ ID NOS: 9703
SOFTWARE: Patentin version 3.0
SEQ ID NO 3026
LENGTH: 17
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Publication No. US20040077565A1
GENERAL INFORMATION:
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Best Local Similarity
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NAME/KEY: misc_feature
ORGANISM: Homo sapiens
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Local Similarity 64.7%;
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; Pred. No. 10;
4; Mismatches
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Pred. No. 1
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Sequence 7674, Application US/10287949A

Publication No. US20040102389A1

GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
ITITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Rel
ITITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
ITITLE OF INVENTION Lovels of Vascular Endothelial Growth Factor Receptor
CURRENT APPLICATION UNMBER: US/10/287,949A
CURRENT APPLICATION UNMBER: US/10/287,949A
CURRENT FILING DATE: 2003-04-11
NUMBER OF SEQ ID NOS: 20082
SOFTWARE: Patentin version 3.0
SEQ ID NO 7674
LENGTH: 17
TYPE: RNA
OPEDATION: Unmo carriers
                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Rel; TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Rel; FILE REFERENCE: MBHB00-876-N (400/049); CURRENT APPLICATION NUMBER: US/10/138,674; CURRENT FILING DATE: 2002-05-03; NUMBER OF SEQ ID NOS SEQ OF NOS SEQ OF SEQ TO NOS SEQ OF SEQ
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                                                                                                                                                                                                           ; TYPE: RNA
; ORGANISM: Homo sapiens
US-10-138-674-5840
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Publication No. US20040077565A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Payco, Pam
                                                                  Query Match
Best Local Similarity
Matches 9; Conserv
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Best Local
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       CTGACTGATGTTGAG 1729
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                                                                         Conservative
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                                                                                                  27.9%; Score 13.4; 60.0%; Pred. No. 10
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Pred. No. 1
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Pred. No. 1
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Sequence 5840, Application US/10287949A
Publication No. US20040102389A1
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: Escobedo, Jaime
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                                                                                             PRIOR APPLICATION NUMBER: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR PILING DATE: 2000-09-27
PRIOR PILING DATE: 2000-09-27
PRIOR PILING DATE: 2001-01-30
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TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00-876-N (400/049)
CURRENT APPLICATION NUMBER: US/10/287,949A
CURRENT FILING DATE: 2003-04-11
NUMBER OF SEQ ID NOS: 20822
SOFTWARE: Patentin version 3.0
SEQ ID NO 5840
LENGTH: 16
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Best Local :
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GENERAL INFORMATI
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CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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TYPE: RNA
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                                                  FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00663
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00662 FILING DATE: 2001-01-30
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Stinchcomb, Daime
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Pred. No. 1
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PRIOR FILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/US01/00670

PRIOR FILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: US 60/234,687

PRIOR APPLICATION NUMBER: US 60/266,860

PRIOR APPLICATION NUMBER: US 60/266,860

PRIOR FILING DATE: 2001-02-05

NUMBER OF SEQ ID NOS: 15752

SOFTWARE: Acomica Sequence Listing Engine
SEQ ID NO 8341

LENGTH: 17
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Matches
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SOFTWARE:
                                                        PRIOR FILING DATE: 2001-01-30 PRIOR APPLICATION NUMBER: US PRIOR FILING DATE: 2000-09-21 PRIOR APPLICATION NUMBER: US PRIOR FILING DATE: 2001-02-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MOSCIE
FILE REFERENCE: ADEMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
                                         NUMBER OF SEQ ID NOS: 15752
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les 13; Conserv
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APPLICATION NUMBER: PCT/US01/00661
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APPLICATION NUMBER: PCT/US01/00662
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APPLICATION NUMBER: PCT/US01/00667
                                                                                                                         APPLICATION NUMBER: US 60/234,687
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FILING DATE: 2001-01-30
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  : Aeomica Sequence Listing Engine
8342
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; LENGTH: 17
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-8342
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US-09-866-108-8343
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US-09-866-108-8343/c
                                  Best Local Similarity 100. Matches 13; Conservative
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LENGTH: 17
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                                                                       Query Match
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Aeomica Sequence
                                                                                                                                                                                                                                        PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
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CURRENT ETLING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
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TILE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
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APPLICATION NUMBER: PCT/US01/00662
FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00668
FILING DATE: 2001-01-30
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16 TGCTGACTGATGT 4
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APPLICATION NUMBER: US 60/234,687
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APPLICATION NUMBER: PCT/US01/00665
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SOFTWARE: Aeomica Sequence
SEQ ID NO 8344
                                                                                  Query Match
Best Local
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APPLICANT: GU, Yizhong
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
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CURRENT FILING DATE: 2001-05-25
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ITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
                             1713 TGCTGACTGATGT 1725
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PENN, Sharron
HANZEL, David
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CHEN, Wenshen
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RESULT 18 US-09-866-108-8345/c

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Sequence 8345, Application US/09866108 Patent No. US20020048800A1 GENERAL INFORMATION:

APPLICANT: GU, Yizhong

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GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals, Inc.

APPLICANT: Pavco, Pam

APPLICANT: McSwiggen, Jim

APPLICANT: Stinchcomb, Dan

APPLICANT: Stinchcomb, Dan

APPLICANT: Stinchcomb, Dan

APPLICANT: Escobedo, Jaime

TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions

TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor

FILE REFERENCE: MBHB00-876-N (400/049)

CURRENT APPLICATION NUMBER: US/10/138,674

CURRENT APPLICATION NUMBER: US/10/138,674

CURRENT FILING DATE: 2002-05-03

NUMBER OF SEQ ID NOS: 20822
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ORGANISM: Homo sapiens
-09-866-108-8345
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PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR APPLICATION NUMBER: PCT/US01/00667
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SOFTWARE: Aeomica Sequence
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CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
FILE REFERENCE: AECMICA-7
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APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
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APPLICATION NUMBER: US 60/266,860
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00661
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APPLICATION NUMBER: PCT/US01/00664
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TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00-876-N (400/049)
CURRENT APPLICATION NUMBER: US/10/287,949A
CURRENT FILING DATE: 2003-04-11
NUMBER OF SEQ ID NOS: 20822
SOTTWARE: Patentin version 3.0
SEQ ID NO 5841
LENGTH: 16
TYPE: RNA
ORGANISM: Homo sapiens
US-10-287-949A-5841
Search completed: July 13, 2004, 11:06:29 Job time : 1 secs
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Best Local
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Publication No. US20040102389A1
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APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
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